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Result
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length: 2000000000
Match
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Gapop 10.0 ,
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1047
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Listing first 45 summaries
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geneseqp2003as:*
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2, 2006, 12:47:32 ; Search time 61 Seconds (without alignments) 1476.601 Million cell updi
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ADM83803
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ADJ70015
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ADM83812
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Biocceleration
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135.5	287	329.5	349	465	465	465	465	584	584	584	584	591	591	591	591	591	606	614	945.5
12.9	27.4	31.5	33.3	44.4	44.4	44.4	44.4	•	55.8	55.8	55.8	56.4	56.4	56.4	56.4	56.4	57.9	58.6	90.3
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Adp26646 Mouse		Abg12341 Nove	Adm83815 Yeast	Adm83805 Humar	Adm83795 Humar	Adm83836 Human	Add84539 1211	Adm83837 Human	Adm83806 Human	Adm83797 Hum	Add84541 121	Adm83844 Human	Adm83807 Humar	Adm83845 Human	Adm83799 Hum	Add84543 121	Aam41829 Human	Adm83843 Human	Adm83852 Human
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ALIGNMENTS

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RESULT 1
ADD84537
 The present invention describes a composition (I) comprising a substance that modulates the status of 121PIF1 (gene and encoded protein), or a molecule that is modulated by 121PIF1, where the status of a cell that expresses 121PIF1 is modulated. The human 121PIF1 gene maps to chromosome 4q. (I) has cytostatic activity, and can be used in gene therapy, and in vaccines. The composition (I) can be used for diagnosing, preventing, prognosticating or treating patients with cancer that expresses 121PIF1, such as breast, colon, ovarian or lung cancer. The 121PIF1 gene or its
                                                                                                                                                                                                Composition comprising a substance that modulates the status of 121P1F1, useful in diagnosing, preventing, prognosticating or treating patients with cancer that expresses 121P1F1, such as breast, colon, ovarian or
                                                                                                                                                    Claim 19; Fig 2A; 285pp; English.
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N-PSDB; ADD84536.
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                                                                                                                                                                                                                                                                                                                                                                                           05-MAR-2001; 2001US-00799250.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene therapy; vaccine; cancer; immune
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fragment can be used to elicit a humoral or cellular
                                                                                                                                                                                   lung cancer.
                                                                                                                                                                                                                                                                                                               Jakobovits A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        121P1F1; 121P1F1 modulation; human; chromosome 4q; cytostatic; gene therapy; vaccine; cancer; immune response; immunisation.
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RESULT 2
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AC ADJ7
AC ADJ7
AC ADJ7
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DE Huma
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W mitc
KW mitc
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KW mice
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mitochondrial; human; screening assay; diabetes mellitus;
Huntington's disease; osteoarthritis;
Leber's hereditary optic neuropathy; LHON;
mitochondrial encephalopathy lactic acidosis and stroke; MELAS;
myoclonic epilepsy ragged red fibre syndrome; MERRF; cancer;
neuroprotective; nootropic; antidiabetic; anticonvulsant; antiarthritic;
                        Identifying a mitochondrial target for drug screening assays and treating diseases associated with altered mitochondrial function, comprises detecting a modified polypeptide in a sample and correl with the disease.
                                                                                                                                                                                                                                                                                                                                                               12-APR-2002; 2002US-0372843P
17-JUN-2002; 2002US-0389987P
20-SEP-2002; 2002US-0412418P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-APR-2003; 2003WO-US010870.
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Claim 1;
                                                                 This invention relates to novel mitochondrial targets that can be
                                                                          SEQ
                                                                          ID NO
                                                                          1821; 180pp;
                                                                          English.
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CC for therapeutic intervention in treating a disease associated with CC altered mitochondrial function. Specifically, it refers to a method for CC identifying proteins of the human heart mitochondrial proteome that are CC useful for drug screening assays, as well as therapeutic targets. The CC present invention describes a method for identifying such proteins that CC can be used in the treatment of various diseases associated with altered CC contendrial function including diabetes mellitus, Huntington's disease, CC osteoarthitis, Leber's hereditary optic neuropathy (LHON), mitochondrial CC encephalopathy lactic acidosis and stroke (MELAS), myoclonic epilepsy CC ragged red fibre syndrome (MERRF) or cancer. Accordingly, these CC compositions have neuroprotective, nootropic, antidiabetic, antiarthritic, osteopathic, ophthalmological and CC cytostatic activities. This polypeptide sequence is a human heart or contended to the invention. used

Sequence A

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ADM63851
ID ADM6
XX ADM6
XX ADM6
XX ADM6
XX ADM6
XX Huma
XX Huma
XX Huma
XX Huma
XX Cerv
XX Cerv
XX US20
XX US20
XX US20
XX O4-E
XX O4
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Best Local
                                                                              (CHAL/)
(HUBE/)
(RAIT/)
(FARI/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human leukocyte
colon cancer; lu
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thes 205;
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GEW.
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                                                                              RAITANO A B. FARIS M.
                                                                                                                                                            CHALLITA-BID P
  JAKOBOVITS
                                                                                                                                     HUBERT R S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene 121P1F1 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene 121P1F1; cytostatic; cancer; chromosome 4q; HLA;
te antigen; prostate cancer; bladder cancer; kidney cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antigen; prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          stomach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer; pancreatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein; 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
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                                                                                                                                                                 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 r; pancreatic cancer; breast cancer; cancer; gene therapy; vaccine.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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CC antibody, a non-human transgenic animal that produces an antibody, a hybridoma that produces an antibody, a single chain monoclonal antibody that immunospecifically binds to a 121P1F1-related protein (comprising CC the variable domains of the heavy and light chains of a monoclonal comprising a polynucleotide that encodes a single chain monoclonal antibody, a vector comprising a polynucleotide that encodes an analogue compile, delivering a cytotoxic agent or a diagnostic agent to a cell chain the expresses 121P1F1, inhibiting growth of cancer cells that expresses 121P1F1 (comprising administering to the cells the composition), treating a patient who bears cancer cells that expresses 121P1F1, inducing an immune cresponse directed to 121P1F1, inducing an immune cresponse directed to 121P1F1, inducing an immune composition may comprise a polynucleotide that immune composition may comprise a polynucleotide in a biological sample from a patient who has or who is suspected of having cancer, monitoring the composition may comprise a polynucleotide that comprises a 121P1F1 cellated protein or polynucleotide in a biological sample from a patient who has or who is suspected of having cancer. The composition may comprises a polynucleotide that comprises a 121P1F1 concer entire amino acid sequence of 121P1F1 having cancer. The composition may comprises a polynucleotide that the coding sequence does not encode the entire amino acid sequence of 121P1F1 (ADM83793. The peptides being HIA (human leukocyte antigen)-binding epitopes from 121P1F1 or its greventing cancer, preferably prostate cancer, bladder cancer, kidney cancer, colon cancer, lung cancer, pancreatic cancer, breast cancer, kidney cancer that expresses or overexpresses concernical cancer or stomach cancer. The composition can also be used as a cervical cancer or stomach cancer that expresses or overexpresses concernical cancer is a 121P1F1 protein (full-length or fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            recombinant protein comprising an antigen-binding region of a monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     composition comprising the novel composition in a human unit dose form,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 5; SEQ ID NO 61; 211pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treating or preventing cancer e.g. prostate, bladder, colon, breast or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New composition comprising a substance that modulates the status of 121P1F1 or a molecule that is modulated by 121P1F1, useful for detecting,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-060522/06
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KRKFGFEENKIDRTFGIPEDFDYID 205
                                                                                           TRLAKELSSLRDQREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWA
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                                                                                                                                                                                                                                                      MSKKKGLSAEEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMV
                                                             TRLAKELSSLRDQREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWA
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                                                                                                                                                                                                                                                                                                                                                 Score 1047;
Pred. No. 1.8
D; Mismatches
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                                                                                                                                                                                                                                                                                                                                               7; DB 8;
1.8e-88;
thes 0;
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                                                                                                                                                                                                                                                                                                                                                                                              Length 205;
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KRKFGFEENKIDRTFGIPEDFDYID

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The invention relates to a composition comprising a substance that commodulates the status of 121P1F1 (a protein encoded by a cancer expressed composition or a molecule that is modulated by 121P1F1 where status of the cell composition comprising the novel composition in a human unit dose form, a crecombinant protein comprising an antipody, an onn-human transgenic animal that produces an antibody, a non-human transgenic animal that produces an antibody, a cariable domains of the heavy and light chains of a monoclonal comprising a polynucleotide that encodes a single chain monoclonal antibody, a vertor comprising a polynucleotide that encodes an analogue peptide, delivering a cycotoxic agent or a diagnostic agent to a cell chain monoclonal antibody, a polynucleotide that encodes an analogue comparising administering growth of cancer cells that expresses 121P1F1, inhibiting growth of cancer cells that expresses 121P1F1 (comprising administering to the cells the composition), treating capanter two bears cancer cells that express 121P1F1, generating a mammalian immune response directed to 121P1F1, inducing an immune capanter to a cell capanter who has or who is suspected of having cancer, monitoring the gresence of cancer in an individual and an assay for detecting the composition may comprise a polynucleotide that comprises a 121P1F1.

CC related protein coding sequence provided that the coding sequence does not encode the entire amino acid sequence of 121P1F1 (ADMS3793. The subside given in 16 Tables (given in the specification), the peptides being HLA (human Telated protein in the specification), the peptides being HLA (human Telated protein in the specification) the peptides being HLA (human Telated protein in the specification).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human leukocyte
colon cancer; lu
cervical cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer.
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HUBERT R S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; cancer gene 121P1F1; cytostatic; cancer; chromosome 4q; HLA; human leukocyte antigen; prostate cancer; bladder cancer; kidney cancer; colon cancer; lung cancer; pancreatic cancer; breast cancer;
                                                             New composition comprising a substance that modulates the status 121P1F1 or a molecule that is modulated by 121P1F1, useful for de treating or preventing cancer e.g. prostate, bladder, colon, breat
                                                                                                                                                                                                                                                            Challita-Eid PM,
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                                          treating or preventing 
lung cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                        (HUBE/)
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) HUBERT R S.
) RAITANO A B.
) FARIS M.
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                                                                                                                                                                                                                                                              Hubert
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Pred. No. 1.8e-88;
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Disclosure;
SEQ ID
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22;
211pp; English
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composition compilising the novel composition in a minimal constitution of recombinant protein comprising an antigen-binding region of a monoclonal contribody, a non-human transgenic animal that produces an antibody, a contribody, a recombinant produces an antibody, a chair monoclonal antibody, a vector comprising a polynucleotide that encodes a single chain monoclonal antibody, a polynucleotide that encodes a single chain monoclonal antibody, a polynucleotide that encodes a single chair monoclonal antibody, a polynucleotide that encodes an analogue comprising administrating to the calls the condess a single craim monoclonal antibody, a polynucleotide that encodes an analogue comparising administrating to the cells that expresses calls that expresses cancer calls that expresses cancer calls that expresses cancer response, monitoring 121PIF1 gene products in a biological sample from a patient who has or who is suspected of having cancer, monitoring the composition may comprise a polynucleotide that comprises a libratering the composition may comprise a polynucleotide that comprises a libratering the composition may comprise a polynucleotide that the coding sequence of cancer in an individual and an assay for detecting the composition may comprise a polynucleotide that comprises a libratering cancer. The composition and in 16 Tables (given in the specification), the peptides cancer, preferably prostate cancer, bladder cancer, kidney cancer, colon cancer, preferably prostate cancer, bladder cancer, kidney cancer, colon cancer, preferably prostate cancer, bladder cancer, beast cancer, cancer for 121PIF1. The gene for 121PIF1 processes or oteracer, breast cancer, cancer for 121PIF1 processes or oteracer as stomach cancer that expresses or overexpresses cancer.

Composition to treat or prevent cancer that expresses or overexpresses cancer. gene) or a molecule that is modulated by 121PIFI where status of the cell that expresses 121PIFI is modulated. Also included are a pharmaceutical composition comprising the novel composition in a human unit dose form, a modulates the status invention relates to a composition comprising a substance that ulates the status of 121P1F1 (a protein encoded by a cancer exp ates the status of 121P1F1 (a proof or a molecule that is modulated is a 121P1F1 protein (full-length or fragment). expressed

Sequence 205 AA;

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Best Local
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                                                                                                                         DCERIGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGRCETEER 120
                                                                                                                                                                              MSKKKGLSABEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMV
                       KRKFGFEENKIDRTFGIPEDFDYID 205
                                                                              TRLAKELSSLRDQREQLKAEVEKYKDCDPQVVEBIRQANKVAKEAANRWTDNIFAIKSWA 180
                                                                                                         DCERIGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGRCETEER
                                                                                                                                                             MSKKKGLSAEEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMV
KRKFGFEENKIDRTFGIPEDFDYID
                                                     TRLAKELSSLRDQREQLKAEVEKYKDCDPQVVEBIRQANKVAKBAANRWTDNIFAIKSWA
                                                                                                                                                                                                                   Conservative
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Pred. No. 1.8e-88;
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ADM83793

standard;

protein; 205

03-JUN-2004 (first entry)

Human cancer gene 121P1F1 protein #1.

RESULT 6
ADM83793
ID ADM8
XX
AC ADM8
XX
DT 03-J
XX
DE Huma
XX
KW Huma Human; cancer gene 121P1F1; cytostatic; cancer; chromosome human leukocyte antigen; prostate cancer; bladder cancer; k bladder cancer; kidney cancer; 4q;

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CC recombinant protein comprising an antispen-binding region of a monoclonal antibody, a non-human transgenic animal that produces an antibody, a chain monoclonal antibody, a chain monoclonal antibody, a chain monoclonal antibody, a chain monoclonal antibody comprising of the heavy and light chains of a monoclonal antibody chain monoclonal antibody, a polymucleotide that encodes a single chain monoclonal antibody, a polymucleotide that encodes an analogue chain monoclonal antibody, a polymucleotide that encodes an analogue chain monoclonal antibody, a polymucleotide that encodes an analogue chain monoclonal antibody, a polymucleotide that encodes an analogue chain monoclonal antibody, a polymucleotide that encodes an analogue chain monoclonal antibody, a polymucleotide that encodes an analogue peptide, delivering a cytotoxic agent or a diagnostic agent to a cell chair expresses 121PIF1, inhibiting growth of cancer cells that expresses cells that expresses analogue composition, treating a commandian immune response directed to 121PIF1, inducing an immune composition and products in a biological sample from a patient who has or who is suspected of having cancer, monitoring the gresence of allers the chair of polymucleotide in a biological sample from a patient who has or who is suspected of having cancer. The composition may comprise a polymucleotide that comprises a 121PIF1 composition of the composition of the polymucleotide in a biological sample from a composition and comprises a polymucleotide that the coding sequence does composition in the specification), the peptides concer, colon cancer, preferably prostate cancer, bladder cancer, kidney cancer, colon cancer, preferably prostate cancer, bladder cancer, breast cancer, colon cancer, prevent cancer that expresses or overexpresses a concer of 121PIF1 is located on chromosome 4q. The present concernance is a 121PIF1 protein (full-length or fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a composition comprising a substance that modulates the status of 121P1F1 (a protein encoded by a cancer expressed gene) or a molecule that is modulated by 121P1F1 where status of the cell that expresses 121P1F1 is modulated. Also included are a pharmaceutical composition comprising the novel composition in a human unit dose form, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 19; SEQ ID NO 3; 211pp; English
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cervical cancer; stomach cancer; gene therapy; vaccine.
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modulates the status of 121PIFI (a protein encoded by a cancer expressed gene) or a molecule that is modulated by 121PIFI where status of the cell that expresses 121PIFI is modulated. Also included are a pharmaceutical composition comprising the novel composition in a human unit dose form, a recombinant protein comprising an antigen-binding region of a monoclonal antibody, a non-human transgenic animal that produces an antibody, a hybridoma that produces an antibody, a single chain monoclonal antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human
                                                                                                                                                  New composition comprising a substance that modulates the status of 121P1F1 or a molecule that is modulated by 121P1F1, useful for detecting, treating or preventing cancer e.g. prostate, bladder, colon, breast or
                                                                                                                                                                                                                                              Challita-Eid PM,
                                                                                                                                                                                                                                                                                                             (HUBE/)
(RAIT/)
(FARI/)
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                                                                                                                  Disclosure; SEQ
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                                                                                                                                                                                                                                   Jakobovits
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                                   28-FEB-2002; 2002US-00087190.
                                                                                                               04-DEC-2003
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Pred. No. 1.8e-88;
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MSKKKGLSAEEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMV

60

Query Match Best Local Matches

Local Similarity 205;

100.0%;

Score 1047; DB 8; Pred. No. 1.8e-88; ; Mismatches 0;

Indels Length

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CC peptide, delivering a cytotoxic agent or a diagnostic agent to a cell CC peptide, delivering a cytotoxic agent or a diagnostic agent to a cell CC that expresses 121PIF1, inhibiting growth of cancer cells that expresses (CI 12PIF1 (comprising administering to the cells the composition), treating CC apatient who bears cancer cells that express 121PIF1, generating a cytotoxic glene products in a biological sample from a CC response, monitoring 12PIF1 generating an immune CC patient who has or who is suspected of having cancer, monitoring the CC presence of cancer in an individual and an assay for detecting the CC presence of a 12PIF1-related protein or polynucleotide in a biological composition may comprise a polynucleotide that comprises a 12PIF1-CC composition may comprise a polynucleotide that conding sequence CC related protein coding sequence provided that the coding sequence does CC not encode the entire amino acid sequence of 12PIF1 (ADM83793). The CC substance also comprises a polynucleotide that encodes at least one CC peptide given in 16 Tables (given in the specification), the peptides being HLA (human leukocyte antigen)-binding epitopes from 12PIF1 or its CC preventing cancer, preferably prostate cancer, bladder cancer, kidney CC cancer, colon cancer, preferably prostate cancer, bladder cancer, kidney CC cancer, color cancer, preferably prostate cancer, breast cancer, kidney CC cancer, to treat or prevent cancer that expresses or overexpresses CC vaccine to treat or prevent cancer that expresses or overexpresses CC cancer as a 12PIFI protein (full-length or fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene) or a molecule that is modulated by 121PIFI where status of the cell that expresses 121PIFI is modulated by 121PIFI where status of the cell that expresses 121PIFI is modulated. Also included are a pharmaceutical composition comprising the novel composition in a human unit dose form, a recombinant protein comprising an antigen-binding region of a monoclonal antibody, a non-human transgenic animal that produces an antibody, a hybridoma that produces an antibody, a single chain monoclonal antibody that immunospecifically binds to a 121PIFI-related protein (comprising the variable domains of the heavy and light chains of a monoclonal antibody), a vector comprising a polynucleotide that encodes a single chain monoclonal antibody, a polynucleotide that encodes an analogue peptide, delivering a cytotoxic agent or a diammostic according.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a composition comprising a substance that modulates the status of 121P1F1 (a protein encoded by a cancer expens) or a molecule that is modulated by 121P1F1 where status of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New composition comprising a substance that modulates the status of 121P1F1 or a molecule that is modulated by 121P1F1, useful for detecting, treating or preventing cancer e.g. prostate, bladder, colon, breast or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-FEB-2001; 2001US-00779250
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                                              is a 121P1F1
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                                              protein (full-length or fragment).
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Daughtery SC, Dam TC,
Peralta CH, David MH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-MAR-2001;
29-MAR-2001;
16-MAY-2001;
17-MAY-2001;
                                                                          The invention relates to a secretory polynucleotide (designated sptm) comprising any of 567 polynucleotide sequences (ABZ35837-ABZ36403), a naturally occurring polynucleotide sequence it least 90 % identical to the polynucleotide sequence, a polynucleotide complementary to them or an RNA equivalent of them. The polypeptide or polynucleotide are useful for treating, preventing or diagnosing a disease or condition associated with the expression of functional SPTM. These are particularly useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    multiple sclerosis; Parkinson's disease; cell proliferative disorder; anti-inflammatory; immunosuppressive; neuroprotective; nootropic; neuroleptic; anticonvulsant; cytostatic; antiparkinsonian; anxiolytic; antipsoriatic; antianaemic; anti-HIV; human immunodeficiency virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; SPTM; autoimmune disorder; inflammatory disorder; AIDS; asthma; Crohn's disease; neurological disorder; epilepsy; cance Huntington's disease; Alzheimer's disease; Creutzfeldt-lakob di
                        diagnosing, treating or preventing autoimmune/inflammatory disorders (e.g. acquired immunodeficiency syndrome, anaemia, asthma or Crohn's
                                                                                                                                                                                                                                                                                                  Claim 27;
                                                                                                                                                                                                                                                                                                                                                                       New human secretory proteins and polynucleotides, useful for diagnosing, treating or preventing autoimmune/inflammatory disorders (e.g. AIDS), neurological disorders (e.g. Alzheimer's), or cell proliferations or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Daffo
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19-JUN-2001;
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                                                                                                                                                                                                                                                                                               SEQ
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     neurological
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2001US-0280068P.

; 2001US-0291280P.

; 2001US-0291829P.

; 2001US-0291849P.

; 2001US-029428P.

; 2001US-0299428P.

; 2001US-0299776P.

; 2001US-0300001P.
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JY, Tuason O,
TF, Nguyen DA,
Nguyen DA,
Nguyen DA,
Nguyen DA,
TP, Urashka ME;
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     (e.g.
                                                                                                                                                                                                                                                                                               Sequence Listing; English
     epilepsy, Huntington's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gietzen D, Chinn J;
O, Yap PE, Amshey SR;
DA, Kleefeld Y, Gerstin
AJ, Panzer SR, Harris B;
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                                                                                                                    (CHAL/)
(HUBE/)
(RAIT/)
(FARI/)
                      WPI; 2004-060522/06
                                                                                                                                                                               08-FEB-2001; 2001US-00779250
                                                                                                                                                                                                                                                                                                   cervical cancer;
                                                                                                                                                                                                                                                                                                             Human; cancer gene 121P1F1; cytostatic; cancer; chromosome 4 human leukocyte antigen; prostate cancer; bladder cancer; ki colon cancer; lung cancer; pancreatic cancer; breast cancer;
                                                                                                                                                                                                                                                                                                                                                               Human cancer
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                                               Jakobovits
                                                          Challita-Eid
                                                                                                                                                                                                      28-FEB-2002; 2002US-00087190
                                                                                                                                                                                                                                                    US2003223997-A1
                                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                         (AFAR/)
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                                                                                                    CHALLITA-EID P N
HUBERT R S.
RAITANO A B.
FARIS M.
FARR D E H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 MSKKKGLSAEEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                 JAKOBOVITS A.
                                                                                             GE W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DCERIGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGRCETEER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KRKFGFEENKIDRTFGIFEDFDYID
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                                                                                                                                                                                                                                                                                                                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TRLAKELSSLRDQREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MSKKKGLSAEEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KRKFGFEENKIDRTFGIPEDFDYID
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                                              P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                              gene
                                                        PM,
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A
                                                                                                                                                                                                                                                                                                                                                                                      (first entry
                                                                                                                                                                                                                                                                                                   ng cancer; pancreatic stomach cancer; gene
                                                          Hubert
                                                                                                                                                                                                                                                                                                                                                              121P1F1 variant protein
                                                                                                                                                                                                                                                                                                                                                                                                                                     protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                                          RS,
                                                                                                                                                                                                                                                                                                                                                                                                                                       206
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 1047; DB 6;
Pred. No. 1.9e-88;
                                                          Raitano
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                  gene therapy; vaccine
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                                                          ₽В,
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                                                            Afar
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                                                          Ge
                                                                                                                                                                                                                                                                                                                            cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              brain,
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composition comprising

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substance that modulates

the

status

of.

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RESULT 11
ADM83835
ID ADM83
XX
AC ADM83
AC ADM83
XX
DT 03-JU
XX
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Composition may remain crainspelic darking the produces an antibody of that immunospecifically binds to a 121PFF1-related protein (comprising the variable domains of the heavy and light chains of a monoclonal antibody the variable domains of the heavy and light chains of a monoclonal control of the variable domains of the heavy and light chains of a monoclonal control of the variable domains of the heavy and light chains of a monoclonal control of the variable domains of the heavy and light chains of a monoclonal control of the variable down of the composition of the control of the composition may comprise a polynucleotide that comprises a lollogical sample from a patient who has or who is suspected of having cancer. The control of the control of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a composition comprising a substance that modulates the status of 121P1F1 (a protein encoded by a cancer expressed gene) or a molecule that is modulated by 121P1F1 where status of the cel
                                             03-JUN-2004
                                                                                         ADM83835;
                                                                                                                                     ADM83835
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  recombinant protein comprising an antigen-binding region of a monoclonal antibody, a non-human transgenic animal that produces an antibody, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                that expresses 121P1F1 is modulated. Also included are a pharmaceutical composition comprising the novel composition in a human unit dose form,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5; SEQ
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                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                      RTRLAKELSSLRDQREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSW
                                                                                                                                                                                                                                                                                                       AKRKFGFEENKIDRTFGIPEDFDYID 205
                                                                                                                                                                                                                                                                                                                                                                                                                                            DCERIGTSNYYWAFPSKALHARKHKLEVLESQQLSEGSQKHASLQKSIEKAKIGRCETEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DCERIGTSNYYWAFPSKALHARKHKLEVLES-QLSEGSQKHASLQKSIEKAKIGRCETEE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MSKKKGLSAEBKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMV
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                                             (first entry)
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                                                                                                                                     protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 8;
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Human; cancer gene 121P1F1; cytostatic; cancer; chromosome 4q; HLA; human leukocyte antigen; prostate cancer; bladder cancer; kidney cancer; colon cancer; lung cancer; pancreatic cancer; breast cancer; cervical cancer; stomach cancer; gene therapy; vaccine.
                                                                                                                                                                                                                            Human cancer gene 121P1F1 variant
                                                                                                                                                                                                                                     protein
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US2003223997-A1.

28-FEB-2002; 2002US-00087190

08-FEB-2001; 2001US-00779250

(RAIT/) (HUBE/) (CHAL/) HUBERT R S.
RAITANO A B. CHALLITA-EID P M.

(AFAR/) (FARI/) AFAR DEH. GEW. FARIS

JAKOBOVITS A.

Challita-Eid PM, Hubert RS, Raitano AB, Faris ĭ Afar ဝူ Σ

WPI; 2004-060522/06

New composition comprising a substance that modulates the status of 121P1F1 or a molecule that is modulated by 121P1F1, useful for detecting, treating or preventing cancer e.g. prostate, bladder, colon, breast 얹

Example 5; SEQ ID NO 45; 211pp; English

CC peptide, delivering a cytotoxic agent or a diagnostic agent to a cell composition and cytotoxic agent or a diagnostic agent to a cell chat expresses 121PIF1 (comprising administering to the cells the composition), treating CC 121PIF1 (comprising administering to the cells the composition), treating CC a patient who bears cancer cells that express 121PIF1, generating a CC mammalian immune response directed to 121PIF1, inducing an immune CC response, monitoring 121PIF1 gene products in a biological sample from a CC response, monitoring 121PIF1 gene products in a biological sample from a CC patient who has or who is suspected of having cancer, monitoring the CC presence of a 121PIF1-related protein or polymucleotide in a biological CC gample from a patient who has or who is suspected of having cancer. The CC composition may comprise a polymucleotide that comprises a 121PIF1-CC related protein coding sequence provided that the coding sequence does CC not encode the entire amino acid sequence of 121PIF1 (ADM83793). The CC substance also comprises a polymucleotide that encodes at least one CC peptide given in 16 Tables (given in the specification), the peptides CC peptide given in 16 Tables (given in the specification), the peptides CC peptide year interest of sequence, bladder cancer, kidney CC cancer, colon cancer, preferably prostate cancer, bladder cancer, kidney CC cancer, colon cancer, preferably prostate cancer, bladder cancer, kidney CC cancer to treat or prevent cancer that expresses or overexpresses a colon cancer or stomach cancer. The composition can also be used as a vaccine to treat or prevent cancer that expresses or overexpresses. recombinant protein comprising an antigen-binding region of a monoclonal antibody, a non-human transgenic animal that produces an antibody, a hybridoma that produces an antibody, a single chain monoclonal antibody that immunospecifically binds to a 121PIF1-related protein (comprising the variable domains of the heavy and light chains of a monoclonal antibody), a vector comprising a polynucleotide that encodes a single chain monoclonal antibody, a polynucleotide that encodes an analogue gene) or a molecule that is modulated by 121PIF1 where status of the cell that expresses 12IPIF1 is modulated by 12IPIF1 where status of the cell composition comprising the novel composition in a human unit dose form, a recombinant protein comprision as assistant in the composition in a human unit dose form, a The invention relates to a composition comprising a substance that modulates the status of 121P1F1 (a protein encoded by a cancer exp a 121P1F1 (full-length or fragment)

SO

Sequence

206

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RESULT 12
ADM83814
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        The invention relates to a composition comprising a substance that modulates the status of 121PIF1 (a protein encoded by a cancer expressed gene) or a molecule that is modulated by 121PIF1 where status of the cell that expresses 121PIF1 is modulated. Also included are a pharmaceutical
                                                             Example 2; SEQ ID NO 24; 211pp;
                                                                                            treating or preventing cancer
                                                                                                                 New composition comprising
                                                                                                                                        WPI; 2004-060522/06.
                                                                                                                                                            Challita-Eid PM,
Jakobovits A;
                                                                                                                                                                                                                                                                                  08-FEB-2001;
                                                                                                                                                                                                                                                                                                      28-FEB-2002; 2002US-00087190.
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                                                                                                                                                                                                                                                                                                                                                 US2003223997-A1
                                                                                                                                                                                                                                                                                                                                                                      Homo
                                                                                                                                                                                                                                                                                                                                                                                           cervical cancer;
                                                                                                                                                                                                                                                                                                                                                                                                      colon
                                                                                                                                                                                                                                                                                                                                                                                                                 human
                                                                                                                                                                                                                                                                                                                                                                                                                           Human;
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(HUBE/)
(RAIT/)
(FARI/)
                                                                                                                                                                                                      (AFAR/)
(GEWW/)
                                                                                                         21P1F1 or a molecule that
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                                                                                                                                                                                                                                                                                                                                                                                                  ; cancer gene 121P1F1; cytostatic; cancer; chromosome 4q; HLA; leukocyte antigen; prostate cancer; bladder cancer; kidney cancer; lung cancer; pancreatic cancer; breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             205;
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                                                                                                                                                                                                      AFAR DEH.
GEW.
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HUBERT R S.
RAITANO A B.
                                                                                                                                                                                            JAKOBOVITS
                                                                                                                                                                                                                             FARIS M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RTRLAKELSSLRDQREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSW 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DCERIGTSNYYWAFPSKALHARKHKLEVLESQELSEGSQKHASLQKSIEKAKIGRCETEE
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                                                                                                                                                                                                                                                                                                                                                                                                                                              gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                           stomach cancer;
                                                                                                                                                                        Hubert
                                                                                                                                                                                                                                                                                                                                                                                                                                                121P1F1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
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                                                                                                                                                                       gg,
                                                                                                                                                                                                                                                                                                                                                                                                                                               protein
                                                                                                       a substance that modulates the status of is modulated by 121PIFI, useful for detecting,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score
Pred.
                                                                                             e.g.
                                                                                                                                                                       Raitano AB,
                                                              English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A
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                                                                                                                                                                                                                                                                                                                                                                                         gene therapy; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                             fragment.
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No. 1.7e-87;
                                                                                             prostate,
                                                                                                                                                                       Faris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 8;
                                                                                             bladder,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
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                                                                                             colon,
                                                                                                                                                                       DEH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1;
                                                                                              breast
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                    cell
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RESULT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC peptide, delivering a cytotoxic agent or a diagnostic agent to a cell
Ct that expresses 121P1F1, inhibiting growth of cancer cells that expresses
CC 121P1F1 (comprising administering to the cells the composition), treating
CC a patient who bears cancer cells that express 121P1F1, generating a
CC mammalian immune response directed to 121P1F1, inducing an immune
CC response, monitoring 121P1F1 gene products in a biological sample from a
CC patient who has or who is suspected of having cancer, monitoring the
CC presence of cancer in an individual and an assay for detecting the
CC presence of a 121P1F1-related protein or polynucleotide in a biological
CC sample from a patient who has or who is suspected of having cancer. The
CC composition may comprise a polynucleotide that comprises a 121P1F1-
CC composition may comprises a polynucleotide that coding sequence does
CC peptide given in 16 Tables (given in the specification), the peptides
CC peptide given in 16 Tables (given in the specification), the peptides
CC peptide given in 16 Tables (given in the specification) the peptides
CC substance also composition is useful for detecting, treating or
CC preventing cancer, preferably prostate cancer, breast cancer, kidney
CC preventing cancer, preferably prostate cancer, breast cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches: 198
                                                                        Human; nootropic; immunosuppressant; cytostatic; gene therapy; cance peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic; chemokinetic; thrombolytic; drug screening; arthritis; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer, colon cancer, lung cancer, pancreatic cancer, breast cancer, cervical cancer or stomach cancer. The composition can also be used as vaccine to treat or prevent cancer that expresses or overexpresses 121P1F1. The gene for 121P1F1 is located on chromosome 4q. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     recombinant protein comprising an antigen-binding region of a monoclonal antibody, a non-human transgenic animal that produces an antibody, a hybridoma that produces an antibody, a single chain monoclonal antibody that immunospecifically binds to a 121P1F1-related protein (comprising the variable domains of the heavy and light chains of a monoclonal antibody), a vector comprising a polynucleotide that encodes a single
                                                                                                                                                                                                         Human polypeptide
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                                                           eukaemia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61
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                                                                                                                                                                                                                                                                                                                                  standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GFEENKIDRTFGIPEDFD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KELSSLRDQREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWAKRKF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGRCETEERTRLA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KGLSAEEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMVDCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      is a 121P1F1 protein (full-length or fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                 GFEENKIDRTFGIPEDFD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KELSSLRDQREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWAKRKF
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Pred. No. 3.6e-85;
0; Mismatches 0;
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WO200153312-A1

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RESULT 14
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Matches 190;
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                                                                                                                                                                                                                                                                                                                                                                                                 immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system disease, such as Alzheimer's, Parkinson's disease, Huntington's disease, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activity caper syndrome. Other uses include the Activin/Inhibin activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and C.N.S disorders. Note: The sequence data for this patent did not form that of the printed specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tang
Wang
Zhou
                           ADD84547
                                                                                                                                                                                                                                                                                                                                                               Sequence 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   encoded polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 4; SEQ ID NO 3188; 10078pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel nucleic acids and polypeptides, useful for treating as central nervous system injuries.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JUL-2000;
03-AUG-2000;
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25-APR-2000;
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                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention relates to human nucleic acids (AAI57798-AAI61369) and oded polypeptides (AAM38642-AAM42213) with nootropic,
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Wang Z,
Goodrich
                                                                                                                                                                                                                                                                        MMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMVDCERIGTSNYYWAFF
                                                                                             GIPEDFDYID
                                                                                                                       GIPEDFDYID 205
                                                                                                                                                  QLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWAKRKFGFEENKIDRTF
                                                                                                                                                                            QLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWAKRKFGFEENKIDRTF
                                                                                                                                                                                                      SKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGRCETEERTRLAKELSSLRDQRE
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                                                                                                                                                                                                                                                        MMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMVDCERIGTSNYYWAFP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; 2000US-00488725.
; 2000US-00552317.
; 2000US-00598042.
; 2000US-00620312.
; 2000US-00623450.
; 2000US-0062191.
; 2000US-00693036.
; 2000US-00727344.
                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                           printed specification
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Wehrman T,
                          protein; 190
                                                                                                                                                                                                                                                                                                                        93.1%; Score 975; DB 4;
100.0%; Pred. No. 7.3e-82;
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hrman T, Xu C,
Drmanac RT;
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Zhang J,
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standard;

GIPEDFDYID

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CC The present invention describes a composition (I) comprising a substance that modulates the status of 121PIF1 (gene and encoded protein), or a CC molecule that is modulated by 121PIF1 (where the status of a cell that CC expresses 121PIF1 is modulated. The human 121PIF1 gene maps to chromosome CC 4q. (I) has cytostatic activity, and can be used in gene therapy, and in CC vaccines. The composition (I) can be used for diagnosing, preventing, CC prognosticating or treating patients with cancer that expresses 12PIF1, CC such as breast, colon, overian or lung cancer. The 121PIF1 gene or its CC fragment can be used to elicit a humoral or cellular immune response. CC 121PIF1 antibodies can be used in active or passive immunisation. 121PIF1 cc polynucleotides are useful as probes and primers for the amplification or CC detection of 121PIF1 genes, as coding sequences for directing the expression of 121PIF1 polypeptides, or as tools for modulating or inhibiting the expression of 121PIF1 genes. The present sequence is used CC in the exemplification of the present invention.
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Best Local Similarity
Matches 190; Conserv
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                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121P1F1; 121P1F1 modulation; human; chromosome 4q; cytostatic; gene therapy; vaccine; cancer; immune response; immunisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 19; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-MAR-2001; 2001US-00799250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-FEB-2002; 2002WO-US006242
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer.
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DB; ADD84546.
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                                                                                                                                                                                                      16 MMBIFSBTKDVFQLKDLBKIAPKEKGITAMSVKEVLQSLVDDGMVDCBRIGTSNYYWAFP
                                                                                                                                                                              1 MMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMVDCERIGTSNYYWAFP
                                                                                                            SKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGRCETEERTRLAKELSSLRDQRE 120
                                                                                                                                                                                                                                                                                                                        190
 GIPEDFDYID 205
                                       QLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWAKRKFGFEENKIDRTF
                                                                QLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWAKRKFGFEENKIDRTF 195
                                                                                                                                  SKALHARKHKLEVLESOLSEGSOKHASLQKSIEKAKIGRCETEERTRLAKELSSLRDQRE 135
                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                  93.1%; Score 975; DB 7; 100.0%; Pred. No. 7.3e-82;
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                                                                                                                                                                                                                                                                                      Length 190;
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RESULT 15
ADM83857
                                                                                                                                     antibody), a vector comprising a polynucleotide that encodes a single chain monoclonal antibody, a polynucleotide that encodes an analogue peptide, delivering a cytotoxic agent or a diagnostic agent to a cell that expresses 121P1F1 (inhibiting growth of cancer cells that expresses 121P1F1 (comprising administering to the cells the composition), treating a patient who bears cancer cells that express 121P1F1, generating a mammallan immune response directed to 121P1F1, inducing an immune response monitoring 121P1F1 gene products in a biological sample from a patient who has or who is suspected of having cancer, monitoring the presence of cancer in an individual and an assay for detecting the
presence of a 121PIFI related protein or polynucleotide in a biological sample from a patient who has or who is suspected of having cancer. The composition may comprise a polynucleotide that comprises a 121PIFI related protein coding sequence provided that the coding sequence does not encode the entire amino acid sequence of 121PIFI (ADM83793. The substance also comprises a polynucleotide that encodes at least one peptide given in 16 Tables (given in the specification), the peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                        modulates the status of 121P1Fi (a protein encoded by a cancer expressed gene) or a molecule that is modulated by 121P1F1 where status of the cell that expresses 121P1F1 is modulated. Also included are a pharmaceutical composition comprising the novel composition in a human unit dose form, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New composition comprising a substance that modulates the status of 121P1F1 or a molecule that is modulated by 121P1F1, useful for detecting, treating or preventing cancer e.g. prostate, bladder, colon, breast or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; cancer gene 121P1F1; cytostatic; cancer; chromosome 4q; HLA; human leukocyte antigen; prostate cancer; bladder cancer; kidney cancer; colon cancer; lung cancer; pancreatic cancer; breast cancer; cervical cancer; stomach cancer; gene therapy; vaccine.
                                                                                                                                                                                                                                                                                                                                 antibody, a non-human transgenic animal that produces an antibody, a hybridoma that produces an antibody, a single chain monoclonal antibody that immunospecifically binds to a 121PIP-related protein (comprising the variable domains of the heavy and light chains of a monoclonal antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                   recombinant protein comprising an antigen-binding region of a monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 5; SEQ ID NO 67; 211pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention relates to a composition comprising a substance that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RAITANO A B. FARIS M. AFAR D E H. GE W.
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HUBERT R S.
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Search completed: February 2, Job time: 63 secs
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 190 AA;
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                                                                                                                                                                                                                                                                                                                             121
                                                                                                                                                                                                                                                                                                                                                                            136 QLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWAKRKFGFEENKIDRTF
                                                                                                                                                            181
                                                                                                                                                                                                                                             196 GIPEDFDYID 205
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                                                                                                                                                            GIPEDFDYID 190
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                                      2006, 12:48:42
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Post-processing: Minimum Match 0%
Maximum Match 100
Listing first 45
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seq length: 2000000000
PIR 80:*
1: pir1:*
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Gapop 10.0 , Gapext 0.5
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1047
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Copyright
                                                                                                                                                                                                                                                                                                                                                                                         2, 2006, 12:50:47; Search time 43 Seconds (without alignments) 458.708 Million cell updates/sec
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                                                                                                                  100%
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

200 200 200 200 200 200 200 200 200 200	Regult No.
349 168:5 135:5 120:5 121:5 117 116:5 111:4 114:4 114:4 114:1 116:1 1 1 1	Score
33.3 16.8 16.8 17.5 17.5 17.5 17.5 17.5 17.5 17.5 17.5	Query Match I
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hypothetical protein F19B15.200 - Arabidopsis thaliana C.Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 09-Jul-2004 C;Accession: T08972 R;Bevan, M.; Hilbert, H.; Braun, M.; Holzer, E.; Brandt, A.; Duesterhoeft, A.; Bancroft, submitted to the Protein Sequence Database, May 1999 A;Reference number: Z16519 A;Accession: T08972

A;Molecule type: DNA

RESULT 2 T08972

44	43	42	41	40	39	38	37	36	35 5	34	33	32	31	30
106.5	107	107	107	107	107	107	107	107	107.5	107.5	107.5	108	109	109
10.2	10.2	10.2	10.2	10.2	10.2	10.2	10.2	10.2	10.3	10.3	10.3	10.3	10.4	10.4
746 1558	955	284	284	284	284	284	281	280	1169	1137	1085	670	676	284
N	N	N	N	N	N	_	N	N	N	N	N	N	N	N
B71603	S24348	A60597	A25825	B27407	A39816	TMRBA	A34787	A22165	A64505	T19414	F96712	F84899	S00084	S19691
myosin ii neavy ch RESA-H3 antigen PF	myosin heavy chain	tropomyosin 2, fib	tropomyosin alpha	tropomyosin alpha	tropomyosin 2, fib	tropomyosin alpha	tropomyosin 1 alph	tropomyosin alpha	P115 homolog - Met	hypothetical prote	hypothetical prote	hypothetical prote	myosin heavy chain	tropomyosin alpha,

ALIGNMENTS

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RESULT 1
T37610
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A;Molecule type: DNA
A;Molecule type: CMN
A;Residues: 1-210 <AUNN
A;Cross-references: UNIPROT:Q09739; UNIPARC:UPI0000139F78; EMBL:Z54096; PIDN:CAA90804.1;
A;Experimental source: strain 972h-; cosmid c13A11
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R;Hunt, S.; Devlin, K.; Churcher, C.M.; Barrell, B.G.; Rajandream, M.A.; Walsh, S.V. submitted to the RMBL Data Library, September 1995
A;Reference number: Z21730
A;Accession: T37610
A;Accession: T37610
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C;Species: O3-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004
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A; Introns: 22/3
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                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 33.3%; Score 349; DB 2; Length 210; Best Local Similarity 41.6%; Pred. No. 3.3e-16; Matches 84; Conservative 36; Mismatches 76; Indels
                                                                                                       123
     183
                                                     181
                                                                                                                                                     122 RLAKELSSLRD-QREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWA 180
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                                                     KRKFGFEENKIDRTFGIPEDFD 202
                                                                                                  QYTLELLHAKESELKLLKTQLSNLNHCNPETFELKNENTKKYMEAANLWTDQIHTLIAFC
                                                                                                                                                                                                                                                                                                             KGLSLAEKRRIEAIFHDSKDFFQLKEVEKLGSK-KQIVLQTVKDVLQSLVDDNIVKTEK 62
                                                                                                                                                                                                                                                                                                                                               KGLSAEEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMVDCER 64
                                                                                                                                                                                                          IGTSNYYWSFPSDAKRSRESVLGSLQAQLDDLKQKSKTLDENISFEKSKRDNEGTENDAN 122
-RDMGADTNQIREYCSIPEDLD 203
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RESULT 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:P53102; UNIPARC:UPI000013B11A; EMBL:X91489; NID:g1143557; R;Bruschi, C.V.; Coglievina, M.; Bertani, I.; Klima, R.; Zaccaria, P.; Delneri, D. submitted to the Protein Sequence Database, May 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Bertani, I.; Coglievina, M.; Zaccaria, P.; Klima, R.; Bruschi, C.V. submitted to the EMBL Data Library, September 1995
A;Description: The sequence analysis of a 7.9 kb DNA fragment from the left
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hypothetical protein YGL183c - yeast (Saccharomyces cerevisiae)

N;Alternate names: hypothetical protein G1604

C;Species: Saccharomyces cerevisiae

C;Date: 23-Feb-1996 #sequence revision 01-Mar-1996 #text_change 09-Jul-2004

C;Accession: S61134; S64200

C;Accession: S61134; S64200
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A;Cross-references: UNIPROT:Q9SZE5; UNIPARC:UPI00000AAA02; EMBL:AL078470; GSPDB:GN00062;
A;Experimental source: cultivar Columbia; BAC clone F19B15
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A; Residues: 1-174 <BRU>
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A;Accession: S64200
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A; Residues: 1-174 <BER>
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A;Accession: S61134
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                                                                                                                                                                                                                                                                                                                                                                                                                                   ;Gene: SGD:MND1
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;Introns: 13/3;
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Best Local :
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Best Local Similarity
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Map position: 7L
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                                                                              120
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                                                                                                                                                                               104 QKSIEKA-KIGRCETEERTRLAKELSSLRDQREQLKAEVEKYKDCDPQVVEEIR-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                76 SKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGRCETEERTRLAKELSSLRDQRE 135
                                                                                                                                                                                                                                                                                                                  46;
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                                                                                                                                                                                                                                   1 MIVKDLVQQMIDEDGVISVEKCGNINIYWCFKNQTLQKLYDSSELIKKKIQEVKCDIATY
                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                         KQELDKTLATGRRKKFTVGQKSYNRBALLEKRKKIQDEIKK-KSNSLQKIESIRWDAAKI
                                                                              QENKQQIRLKKVHLEKTTDNIEILIDYLYKKFFLKPEQIRKEFGIPEEF 168
                                                                                                                  QANK----VAKEAANRWTDNIFAIKSWAKKFGFEENKIDRTFGIPEDF 201
                                                                                                                                                                                                                                                                        MSVKEVLQSLVD-DGMVDCERIGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASL 103
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27.2%;
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%; Pred. No. 1.9e-06;
23; Mismatches 37
                                                                                                                                                                                                                                                                                                                  42;
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; Pred. No. 0.00
42; Mismatches
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C;Species: Gallus gallus (chicken)
C;Date: 28-Aug-1992 #sequence_revision 28-Aug-1992 #text_change 09-Jul-2004
C;Accession: A41642
R;Gill, S.R.; Schroer, T.A.; Szilak, I.; Steuer, E.R.; Sheetz, M.P.; Cleveland, D.W.
J. Cell Biol. 115, 1639-1650, 1991
A;Title: Dynactin, a conserved, ubiquitously expressed component of an activator of vesi A;Reference number: A41642; MUID:92098576; PMID:1836789
A;Accession: A41642
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N;Alternate names: p150 Glued
C;Species: Mus musculus (house mouse)
C;Date: 28-May-1997 #sequence_revision 18-Jul-1997 #text_change 09-Jul-2004
C;Accession: JC5368
R;Jang, W.; Weber, J.S.; Tokito, M.K.; Holzbaur, B.L.F.; Meisler, M.H.
Biochem. Biophys. Res. Commun. 231, 344-347, 1997
Biochem. Biophys. Res. Commun. 231, 344-347, 1997
A;Title: Mouse p150Glued (dynactin 1) cDNA sequence and evaluation as a candidate
A;Reference number: JC5368; MUID:97223454; PMID:9070275
A;Accession: JC5368
                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-1053 <GIL>
A;Cross-references: UNIPROT:P35458; UNIPARC:UPI000017C003; GB:X62773; NID:g63920; PID:g6
C;Keywords: cytoskeleton
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A41642
dynactin - chicken
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A;Residues: 1-1281 <JAN>
A;Cross-references: UNIPROT:O08788; UNIPARC:UPI0000027AAD; GB:U60312; NID:g2104494; PIDN
A;Experimental source: brain
C;Comment: This protein is a member of the oligomeric dynactin complex that is required
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C; Genetics:
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A;Map position: 6
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        279 ARKEAKEALEAKERYMEEMADTADAIEMATLDKEMAEERAESLQQEVEALKERVDELTTD 338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                62 CERI-----GTSNYYWAFPSKALHAR-KHKLEVLESQLSEGSQKHASLQKSIEKA 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       50;
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                                                                                                                                                                                                                       AKKEAKDALEAKERYMEEMADTADAIEMATLDKEMABERABSLQQEVDSLKEKVEYLTMD 160
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                                                                                                              LEILKHEIEEKGSDGAASSYQVKQLEEQNARLKEALVRMRDLSASEKQEHVKLQKQMEKK
                                                                                                                                                                GMV---DCERIGTSNYYWAFPSKALHARKHKLE---VLESQLSEG-SQKHASLQKSIEKA 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LRETVGDLEAMNEMNDXLQENARETELELREQLD
                                                    KIGRCETEERTRLAKELSSLRDQREQLKASVEK-YKDCD------PQVVEEIRQ 157
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-TELESLROOREKLOEEVKOAEKTVDELKEOVDAALGAEEMVETLTE
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Pred. No. 0.44;
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Pred. No. 0.25;
3; Mismatches 106;
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158

ANKVAKEAANRWTDNIFAIKSWAKRKFGFEENKIDRTFGIPEDFD

202

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hypothetical protein At2333300 [imported] - Arabidopsis thaliana N;Alternate names: hypothetical protein T16824.6 C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 05-Mar-1999 #sequence_revision 05-Mar-1999 #text_change 09-Jul-2004 C;Accession: T02572; F84815 R;Rounsley, S. D.; Kaul, S.; Lin, X.; Ketchum, K.A.; Crosby, M.L.; Brandon, R.(submitted to the EMBL Data Library, August 1998 A;Description: Arabidopsis thaliana chromosome II BAC T16B24 genomic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-1356 <KRUy
A;Cross-references: UNIPROT:Q86UP2; UNIPARC:UPI00000702A8; EMBL:Z22551
A;Cross-references: UNIPROT:Q86UP2; UNIPARC:UPI00000702A8; EMBL:Z22551
A;Cross-references: UNIPROT:Q86UP2; UNIPARC:UPI00000702A8; EMBL:Z22551
A;Cross-references: UNIPROT:Q86UP2; UNIPARC:UPI00000702A8; EMBL:Z22551
A;Title: Molecular cloning and characterization of human kinectin.
A;Reference number: 137947; MUID:95306853; PMID:7787243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #
C;Accession: S32763; I37947
C;Accession: S32763; I37947
R;Kruppa, G.; Fuetterer, A.; Lemke, H.; Kroenke, M.
submitted to the EMBL Data Library, April 1993
A;Description: Cloning and characterization of TAF,
A;Residues: 1-768 <ROU>
A;Residues: 1-768 <ROU>
A;Residues: 1-768 <ROU>
A;Cross-references: UNIPROT:080951; UNIPARC:UPI00000A1172; EMBL:AC004697; NID:g3402671;
A;Experimental source: cultivar Columbia
A;Experimental source: cultivar Columbia
A;Experimental source: cultivar Columbia
A;Exin, X.; Kaul, S.; Rounsley, S.D.; Shea, T.P.; Benito, M.I.; Town, C.D.; Fujii, C.Y.;
R;Lin, X.; Kaul, S.; Rounsley, S.D.; Shea, M.; Vanaken, S.E.; Umayam, L.; Tallon, L.
M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; Vanaken, S.E.; Umayam, L.; Tallon, L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 7
T02572
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                                                                                                                                                        A;Status: translated from GB/EMBL/DDBJ
A;Golecule type: DNA
                                                                                                                                                                                                                 A;Reference number: Z14679
A;Accession: T02572
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Best Local
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Pred. No. 2.2;
31; Mismatches
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                               C.Y.;
lon, L.
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centrosome associated protein CEP250 - human
C;Species: Homo sapiens (man)
C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 21-Jul-2000
C;Accession: T08621
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Molecule type: DNA
A;Residues: 1-768 <STO>
A;Cross-references: UNIPARC:UPI00000A1172; GB:AE002093; NID:g3402677; PIDN:AAC28980.1; C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana A;Reference number: A84420; MUID:20083487; PMID:10617197 A;Accession: F84815
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A;Title: Autoantibodies to a group of centrosomal proteins A;Reference number: Z16462; MUID:98165428; PMID:9506584
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                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: mRNA
A; Residues: 1-2442 < MAC>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         euss, D.; Nierman, W.C.; White, O.; Eisen, J.A.; Salzberg,
                                                                                                                                                                                                                                                                          Matches
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1821
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ALQQEQQQAQGQEERVKEKADALQGALEQAHMTLKERHGELQDH----KEQARR 1870
                              SLRDQREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWAKR
                                                                ---SILLSQREQEIVVLQQQLQEAREQGELKEQSLQSQLDEAQRALAQRDQ-----ELE
                                                                                                   AFPSKALHARKHKLEVLESQLSE----GSQKHASLQKSIEKAKIGRCETEERTRLAKELS
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22.1%;
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                                                                                                                                                                                                                                                                          Score 116.5;
Pred. No. 9;
47; Mismatches
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Pred. No. 2.4;
46; Mismatches
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                                                                                                                                                                                                                                                                                                                                                           R;Whitby, F.G.; Kent, H.M.; Stewart, F.; Stewart, M.; Xie, X.; Hatch, submitted to the EMBL Data Library, April 1992
A;Description: Structure of tropomyosin at 9 Angstroms resolution.
A;Reference number: S24972
A;Accession: S24972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tropomyosin alpha, cardiac - pig
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 20-reb-1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004
C;Accession: S24972
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C;Superfamily: Archaeoglobus fulgidus conserved hypothetical protein
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A;Cross-references: UNIPROT:Q9UZC8; UNIPARC:UPI0000034523; GB:AJ248286; GB:AL096836;
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R;anonymous, Genoscope
submitted to the EMBL Data Library, July 1999
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C;Species: Pyrococcus abyssi
C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                              A; Molecule type: mRNA
A; Residues: 1-284 < WHI>
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                                                                                                                                                                         Similarity
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                                                                                                        MSKKKGLSAE---EKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDD 57
LAEKKATDAEADVASLNRRIQLFEEELDRAQ----ERLA-----TALQKLEEAEKAADE
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                                                                                                                                                   Conservative
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24.1%; Pred. No. 3.5;
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                                                                                                                                                   37;
                                                                                                                                                                       Score 114; DB Pred. No. 1.3;
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  A;Cross-references: UNIPARC:UPI0000137254; EMBL:X86470; NID:g791101; PIDN:CAA60179.1; R;Poehlmann, R.; Philippsen, P.
                                           A; Molecule type: DNA
A; Residues: 1-199 < POE>
                                                                                      A; Reference number: S53896
A; Accession: S53899
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RESULT 12
A32183
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hypothetical protein F10M6.170 - Arabidopsis thaliana (;Species: Arabidopsis thaliana (mouse-ear cress) C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 09-Jul-2004 C;Accession: T05409
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T05409
                                                                                                                                                                                                              tropomyosin TPM1 - yeast (Saccharomyces cerevisiae)
N;Alternate names: protein N2332; protein YNL079c
C;Species: Saccharomyces cerevisiae
C;Date: 08-Sep-1989 #sequence revision 08-Sep-1989 #text_change
C;Date: 08-Sep-1989; S53899; S63011; S63018; S63928
C;Accession: A32183; S53899; S63011; S63018; S63928
R;Liu, H.; Bretscher, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R; Bevan, M.; Weichselgartner, M.; Fartmann, B.; Granderath, submitted to the Protein Sequence Database, February 1998
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A;Cross-references: UNIPROT:P17536; UNIPARC:UPI0000137254; R;Poehlmann, R.; Philippsen, P. submitted to the EMBL Data Library, April 1995
                                                                                                                                           Cell 57, 233-242, 1989
A;Title: Disruption of the single tropomyosin gene in yeast results in the disappearance A;Reference number: A32183; MUID:89195234; PMID:2649250
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A; Note: F10M6.1
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                                                                       A; Molecule type: DNA
A; Residues: 1-199 <LIU>
                                                                                                                     A; Accession: A32183
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A;Accession: T05409
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ----AEFAERSVTKLEKSID
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                                                      EMBL: M25501;
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A;Gene: ATmS-1
C;Superfamily:
C;Kgywords: act
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A;Cross-references: UNIPARC:UPI0000137254; EMBL:Z71355; NID:g1301970; PIDN:CAA95953.1; A;Cross-references: UNIPARC:UPI0000137254; EMBL:Z71355; NID:g1301970; PIDN:CAA95953.1; A;Experimental source: strain S288C
R;Soler-Mira, A.; Saiz, J.E.; Ballesta, J.P.G.; Remacha, M. submitted to the Protein Sequence Database, April 1996
                                                                                                                                                                                R;Luque, E.A.; Spinner, B.J.; Dube, S.; Dube, D.K.; Lemanski, L.F. Gene 185, 175-180, 1997
A;Title: Differential expression of a novel isoform of alpha-tropomyosin in A;Reference number: JC66.98; MUID:97208870; PMID:9055812
A;Contents: skeltal muscle
                                                                                                                                                                                                                                                                                     alpha-tropomyosin S-1 - axolotl
C;Species: Ambystoma mexicanum (axolotl)
C;Date: 11-Apr-1997 #sequence_revision 09-May-1997 #text_change 09-Jul-2004
C;Accession: JC6199
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C; Superfamily: slime mold ribozyme I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Status: nucleic acid sequence not shown;
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A;Title: Sequencing a cosmid clone of Saccharomyces cerevisiae chromosome XIV reveals
A;Reference number: 863925; MUID:96267764; PMID:8701611
A;Accession: 863928
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A; Accession: S63018
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A;Accession: S63011
                                                                               A; Cross-references: UNIPROT: P87349; UNIPARC: UPI00000FB5F3; GB: U33450; C; Comment: This protein is a actin-binding protein.
                                                                                                                        A;Accession: JC6199
A;Molecule type: mRNA
A;Residues: 1-284 <LUQ>
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A;Cross-references: UNIPARC:UPI0000137254;
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                                                                ;Genetics:
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Superfamily: tropomyosin
Keywords: actin binding
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    tapeworm
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R;Clayton, L.; Reinach, F.C.; Chumbley, G.M. Mol. Biol. 201, 507-515, 1988
A;Title: Organization of the hTM(nm) gene. A;Reference number: S02554; MUID:88332987; A;Accession: S06210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Reinach, F.C.; MacLeod, A.R.
Nature 322, 648-650, 1986
A;Title: Tissue-specific expression of the A;Reference number: A24199; MUID:86311274; A;Accession: A24199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Status: not compared with conceptual translation A;Molecule type: DNA A;Residues: 1-285 <CLA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Species: Homo sapiens (man)
C;Date: 02-Jun-1988 #sequence_revision 02-Jun-1988 #text_change
C;Accession: S06210; A24199
                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Superfamily: tropomyosin C;Keywords: alternative splicing; coiled coil; muscle; skeletal muscle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: UNIPARC:UPI000013D90D; GB:X04201; NID:g37429; PIDN:CAA27798.
A;Note: an intronless pseudogene resembling this mRNA is also known
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: mRNA
A; Residues: 1-285 < REI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: UNIPARC:UPI000013D90D R;Reinach, F.C.; MacLeod, A.R.
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                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    162 AKEAANRWTDNIFAIKSWAKRKFGFEENKID 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           133 R----ALKDEEKMELQEI-------QLQEAKHIAEEADRKYEEVARKLVIIEGDL
                                                                                                                                                                             134 R----ALKDEEKMELQEI-------QLKEAKHIAEEADRKYEEVARKLVIIEGDL
                                                                                                                                                                                                                     48 KEVLOSLVDDGMVDCERIGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQKSI 107
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RM MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Altachen R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altachen R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altachen R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., McGun T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Sarchen B.J., Lu X., Gibbs R.A.,
RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA William D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Hakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
Butterfield Y.S.N., Krzywinski M.J., Skalska U., Smailus D.E.,
RA Rodriguez CDNA sequences ";
Brock Mathala and pais of more than 15,000 full-length human and mouse cDNA sequences ";
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Best Local Similarity
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                                                          EMBL; BC027741; AAH27741.1; -; MANA.
Ensembl; ENSMUSG00000033752; Mus musculus
MGI; MGI:1924165; 2610034E18Rik.
InterPro; IPR005647; Mnd1.
                                                                                                                                                                                                                                                              STRAIN=CZECH II;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna; Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                    Submitted (APR-2002) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                          TISSUE=Mammary tumor metastatized Director MGC Project;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mus musculus (Mouse).
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         SEQUENCE
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Pred. No. 8e-57;
         122C3FA9E4325120 CRC64;
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RA Kawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,
RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Airawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Airawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Airawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Kadota K., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Kadota K., Matsuda H.A., Sahburner M., Batalov S., Cassavant T.,
RA Kadota K., Matsuda H.A., Sahburner M., Batalov S., Cassavant T.,
RA Kadota K., Matsuda H.A., Sahburner M., Wagner L., Washio T.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Sakai K., Okido T., Fruno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kaniya M., Lee N.H.,
RA Lyons P., Marchionni L., Mashima J., Mazzarelli J., Bakamoto N.,
RA Marchioli H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Marchioli P., Cannoni A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
Wanshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
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01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Mus musculus 10 days embryo whole body cDNA, RIKEN full-length
enriched library, clone:2610034E18 product:GAJ homolog.
Name=2610034E18Rik,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=C57BL/6J; TISSUE=Whole body;
MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
Carninci P., Hayashizaki Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Mammalia; Eutheria; Buarchontoglires; Glires; Rodentia;
Muroidea; Muridae; Murinae; Mus.
                                          STRAIN=C57BL/6J; TISSUE=Whole body; The FANTOM Consortium,
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                                                                                NUCLEOTIDE SEQUENCE
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RIKEN Genome Exploration Research
alysis of the mouse transcriptome l
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303:19-44(1999).
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89.8%;
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Pred. No.
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    Phase I & II
on functional
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                       Team;
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RESULT 4
Q5FV93 XENTR
ID Q5FV93, XENTR PRELIMINARY;
AC Q5FV93;
DT 10-MAY-2005 (TrEMBLrel. 30,
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RA Adachi J., Aizawa K., Akahira S., Akimura T., Arai A., Aono H.,

RA Adachi J., Aizawa K., Akahira S., Fukunishi Y., Furuno M.,

RA Arakawa T., Bono H., Carninci P., Fukuda S., Fukunishi Y., Furuno M.,

RA Arakawa T., Bono H., Carninci P., Fukuda S., Fukunishi Y., Furuno M.,

RA Imotani K., Ishii Y., Itoh M., Izawa M., Kasukawa T., Kato H.,

RA Kawai J., Kojima Y., Konno H., Kouda M., Koya S., Kurihara C.,

RA Kawai J., Miyazaki A., Nishi K., Nomura K., Numazaki R., Ohno M.,

RA Cazaki Y., Okido T., Owa C., Saito H., Saito R., Sakai C., Sakai K.,

Sano H., Sasaki D., Shibata K., Shibata Y., Shinagawa A., Shiraki T.,

RA Sano H., Sasaki D., Shibata K., Shibata Y., Shinagawa A., Shiraki T.,

RA Sano H., Sazuki H., Tagami M., Tagawa A., Taahaka T.,

RA Sano Y., Toya T., Yamamura T., Yasunishi A., Yoshida K., Yoshino M.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Taahaka T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Taahaka T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Taahaka T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Taahaka T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Yoshida K., Yoshino M.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Takahashi F., Tanaka T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Takahashi F., Tanaka T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Takahashi F., Tanaka T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Takahashi F., Tanaka T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Takahashi F., Tanaka T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Takahashi F., Tanaka T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Shiraki T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Shiraki T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Shiraki T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Shiraki T.,

RA MITEMBARE Y., Suzuki H., Tagami M., Tagawa A., Shiraki T.,

RA MITEMBARE Y., Suzuki H., MITEMBARE Y., Shiraki T.,

RA MITEMBARE Y., Suzuk
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Best Local
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MEDLINE=20530913; PubMed=11076861; DOI=10.1101/gr.152600;

MEDLINE=20530913; PubMed=11076861; DOI=10.1101/gr.152600;

Shibata K., Itoh M., Aizawa K., Nagaoka S., Sasaki N., Carninci P.,

Konno H., Akiyama J., Nishi K., Kitsunai T., Tashiro H., Itoh M.,

Yamamoto R., Matsumoto H., Sakaguchi S., Ikegami T., Kashiwagi K.,

Yujiwake S., Inoue K., Togawa Y., Izawa M., Ohara E., Watahiki M.,

Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,

Yoneda Y., Muramateu M., Inoue Y., Kira A., Hayashizaki Y.;

"RIKEN integrated sequence analysis (RISA) system-384-format

sequencing pipeline with 384 multicapillary sequencer.";

Genome Res. 10:1757-1771(2000).
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MEDLINB-2049974; PubMed-11042159; DOI-10.1101/gr.145100;
MEDLINB-2049974; PubMed-11042159; DOI-10.1101/gr.145100;
Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K.,
Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.;
"Normalization and subtraction of cap-trapper-selected cDNAs to prepare full-length cDNA libraries for rapid discovery of new Genome Res. 10:1617-1630(2000).
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420:563-573(2002).
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89.3%; Pred. No. 9.6e
Live 10; Mismatches
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RESULT
Q5XGY9
ID Q7
AC Q1
DT 21
DT 22
DT 22
DT 21

Q5XGY9; Q5XGY9_XENLA

PRELIMINARY;

205 Å

SENLA

25-OCT-2004 (TremBLrel. 25-OCT-2004 (TremBLrel. 25-OCT-2004 (TremBLrel. LOC495113 protein.

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sequence update)
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RA Altschul S.F., Zeeberg B., Buetcow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetcow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetcow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsish F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Wiltalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Wilting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J.M., Smailus D.E.,
RA Rodriguez A.C., Grimwood J., Schmutz J.M., Smailus D.E.,
RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
RT and mouse cDNA sequences.";
RT proc. Nafl Asad Sci II S.A 90-16404-16001/20021
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Best Local
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SEOUENCE 205 AA; 24130 MW;
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Submitted (FEB-2005) to the
EMBL; BC090139; AAH90139.1;
InterPro; IPR005647; Mnd1.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
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10-MAY-2005 (TrEMBLrel.
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Name=MGC97859;
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181
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                                                                                           TRLAKELSSLRDQREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWA
                                                                                                                                                                     DCERIGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGRCETEER
                                                                                                                                                  DSERIGTSNYFWAFPSKALHARKRKLETLESQFTEVKQKRENLQQSVDKARVGRQDTEER
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KKKFGFEERQIDKNFGIPEDFDYID
                 KRKFGFEENKIDRTFGIPEDFDYID
                                                                        SKLABELASRRHRKEELCAELEKYKECDPDVIEEIROSNKVAKDAVNRWTDNIFAVKSWA
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                                                                                                                                                                                                                                                                                                    81.0%; Score 848; DB 2;
77.1%; Pred. No. 1.3e-44;
tive 29; Mismatches 18
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-; mRNA.
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RA Klausher R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Klausher R.D., Collins F.S., Wagner L.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Butow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Lapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Uddin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Loquellano N.A., Peters G.J., Malek J.A., Gunaratne P.H.,
RA Robards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Mozny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Mozny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Rahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Rahey J., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rahesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length human
"Generation and initial analysis of more than 15,000 full-length human
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Best Local
   ILT 6
106_TETNG
Q4SS06_TETNG
Q4SS06;
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MEDLINE=22341132; PubMed=12454917; DOI=10.1002/dvdy.10174;
Klein S.L., Strausberg R.L., Wagner L., Pontius J., Clifto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Klein S., Gerhard D.S.;
Submitted (OCT-2004) to the
EMBL; BC084288; AAH84288 1;
InterPro; IPR005647; Mnd1.
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Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
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                                                                                                                                                                                                KKKFGFEERQIDKNFGIPEDFDYID
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Pred. No. 1.3e-
27; Mismatches
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                                       PRT;
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RESULT
Q6DC61
ID Q6
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A Mauceli E., Bouneau L., Fischer C., Ozouf-Costaz C., Bernot A.,

A Micaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,

A Nicaud S., Jaffe D., Fisher S., Lutfalla G., Costellano S.,

A Dasilva C., Salanoubat M., Levy M., Boudet N., Castellano S.,

A Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,

A Anthouard C., Skalli Z., Cattolico L., Poulain J., De Berardinis V.,

B Siemont C., Skalli Z., Cattolico L., Poulain J.P., Gouzy J.,

Cruaud C., Duprat S., Brottier P., Coutanceau J.P., Gouzy J.,

A Parra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,

A Parra G., Larder G., Chapple C., McKernan K.J., McEwan P., Bosak S.,

A Kellis M., Volff JN., Guigo R., Zody M.C., Mesirov J.,

A Kellis M., Volff JN., Guigo R., Zody M.C., Mesirov J.,

Laudet V., Schachter V., Outier F., Saurin W., Scarpelli C.,

Wincker P., Lander B.S., Weissenbach J., Roest Crollius H.,

"Genome duplication in the teleost fish Tetraodon nigroviridis reveals

"Genome duplication in the teleost fish Tetraodon nigroviridis reveals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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Best Local
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13-SEP-2005 (T
13-SEP-2005 (T
Chromosome 18
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom
Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleost
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
Tetradontoidea; Tetraodontidae; Tetraodon.
                                                                                                                                                                                                                             QGDC61_BRARE
QGDC61;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genoscope; Whitehead Institute Centre for Genome Research; Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases.

-I- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which
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Nature 431:946-957(2004).
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                                                                                                                                                                                                                                                                                                 BRARE
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(TrEMBLrel. 31, Last sequence update)
(TrEMBLrel. 31, Last annotation updat
(TrEMBLrel. 31, Ast annotation seq
18 SCAF14485, whole genome shotgun seq
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Neoteleostei;
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25-OCT-2004 (TrEMBLre 25-OCT-2004 (TrEMBLre 25-OCT-2004 (TrEMBLre 25-OCT-2007) 2gc:101017 ORFNames=zgc:101017; Brachydanio rerio (Ze

(Zebrafish)

Craniata; (Danio

Vertebrata;

Euteleostomi;

rerio)

(TrEMBLrel. 28, (TrEMBLrel. 28, (TrEMBLrel. 28,

Created)
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Last annotation update)

PRELIMINARY;

PRT;

220

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RESULT 8
Q86EZ8 SCHJA
ID Q86EZ8;
AC Q86EZ8;
DT 01-JUN-2
DT 01-JUN-2
DT 01-OCT-2
DE Clone Zos
OS Schistos
OC Eukaryot
OC Schistos
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RN [1]
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RA Wang Z.J
RA Wang Z.J
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RM MEDLINE-22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,

RA Blatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hullyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hullyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hullyk S.W.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Rodrigues S., Sanchez A.,

RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Willalon D.K., Toynung A.C., Shevchenko Y., Bouffard G.G.,

RA Hakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,

RA Balakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,

RA Balakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,

RA Botherch A., Schein J.E., Jones S.J.M., Warra M.A.,

"Generation and initial analysis of more than 15,000 full-length human

RT Tock. Watl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
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Best Local Similarity
                                                                                                                                                                                                                                       QB6EZ8 SCHJA PRELIMINARY; PRT; 1
Q86EZ8;
Q1-JUN-2003 (TEMBLrel. 24, Created)
Q1-JUN-2003 (TEMBLrel. 24, Last sequer
Q1-OCT-2003 (TEMBLrel. 25, Last annotation of the ZZD1259 mRNA sequence.
Schistosoma japonicum (Blood fluke).
Eukaryota; Metazoa; Platyhelminthes; Tischistosomatidae; Schistosomatidae; Schistosomatidae;
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Pfam; PF03962; Mnd1;
Pfam; PF03962; And1;
NUCLEOTIDE SEQUENCE.

MEDLINB=22879925; Pubmed=12973349; DOI=10.1038/ng1236;

MEDLINB=22879925; Pubmed=12973349; DOI=10.1038/ng1236;

Hu W., Yan Q., Shen D.K., Liu F., Zhu Z.D., Song H.D., Xu X.R.,

Wang Z.J., Rong Y.P., Zeng L.C., Wu J., Zhang X., Wang J.J., Xu

Wang S.Y., Fu G., Zhang X.L., Wang Z.Q., Brindley P.J., McManus
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                                                                                                                                                                                                           NCBI_TaxID=6182;
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Director MGC Project;
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Cyprinidae; Danio.
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Pred. No. 4.8e-32;
0; Mismatches 29
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Schistosoma
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RA Amedeo P., Roncaglia P., Berriman M., Hirt R.P., Mann B.J., Nozaki T.,
RA Amedeo P., Roncaglia P., Berriman M., Hirt R.P., Mann B.J., Nozaki T.,
RA Suh B., Pop M., Duchene M., Ackers J., Tannich E., Leippe M.,
RA Hofer M., Bruchhaus I., Willhoeft U., Bhattacharya A.,
RA Chillingworth T., Churcher C., Hance Z., Harris B., Harris D.,
RA Chillingworth T., Churcher C., Hance Z., Harris B., Harris D.,
RA Guail M.A., Rabbinowitsch B., Norbertczak H., Price C., Wang Z.,
RA Guillen N., Gilchrist C., Stroup S.E., Bhattacharya S., Lohia A.,
RA Guillen N., Gilchrist C., Stroup S.E., Bhattacharya S., Lohia A.,
RA Foster P.G., Sicheritz-Ponten T., Weber C., Singh U., Mukherjee C.,
RA Foster P.G., Sicheritz-Ponten T., Weber C., Singh U., Mukherjee C.,
RA Fraser C.M., Petri W.A., Clark C.G., Embley T.M., Barrell B.,
RFraser C.M., Hall N.,
T'The genome of the protist parasite Entamoeba histolytica.";
Nature 433:865-868 (2005).
C. PARIJON: The sequence shown here is derived from an
EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
                          Query Match
Best Local (
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EMBL; AAFB01000572; EAL45733.1; -;
InterPro; IPR005647; Mnd1.
Pfam; PF03962; Mnd1; 1.
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Q50VW2_ENTHI PRELIMINARY;
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"Evolutionary and biomedical implications complementary DNA resource.";

15.139-147(2003).
                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PubMed=15729342; DOI=10.1038/nature03291;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=HM-1: IMSS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Entamoeba histolytica HM-1:IMSS
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EMBL; AY223066; AAP06089.1; -; mRNA
                                                                                                                                         Hypothetical
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47.4%; Pred. No. 6.4e-21;
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  53;
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Score 459; DB
Pred. No. 1.4e:
53; Mismatches
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                                                                                                                  63AD3885CF0387B3 CRC64;
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Best Local
Matches
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EMBL; AK117713; BAC42364.1; -; mRNA.

EMBL; BT005435; AA063855.1; -; mRNA.
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Seki M., Iida K., Satou M., Sakurai T., Ak
Seki M., Iida K., Satou M., Sakurai T., Ak
Nakajima M., Enju A., Kamiya A., Narusaka
Hayashizaki Y., Shinozaki K.;
Submitted (NOV-2002) to the EMBL/GenBank/I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro; IPR005647; Mnd1.
Pfam; PF03962; Mnd1; 1.
Hypothetical protein.
SEQUENCE 230 AA; 26402 M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Name=At4g29170/F19B15_200; Synonyms=At4g29170;
Arabidopsis thallana (Mouse-ear cress).
Bukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; rosids;
eurosids II, Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAR-2003 (TrEMBLrel. 23, Created)
01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-FEB-2005 (TrEMBLrel. 29, Last annotation update)
Hypothetical protein At4g29170/F19B15_200 (Hypothetical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                QBGYD2_ARATH
QBGYD2;
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                                                                                                                                                                                                                                                               AKDKIGISIYFWSLPSCAGNQLRSVRQKLESDLQGSNKRLAELVDQCEALKKGREESEER
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                                     SNNFPQAKEQLEHLYTEAGITEDFDYIE
                                                                                            KRKFGFEENKIDRTF--
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                                                               -GIPEDFDYID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 440; DB
Pred. No. 1.7e
43; Mismatches
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ka M., Carninci P.,
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O54E86 DICDI PRELIMINARY;
O54E86;
13-SEP-2005 (TrEMBLrel. 31, C
13-SEP-2005 (TrEMBLrel. 31, L
13-SEP-2005 (TrEMBLrel. 31, L
13-SEP-2005 (TrEMBLrel. 31, L
ORFNameseDB0184048;
ORFNameseDB0184048;
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32 ORYSA
06H432 ORYSA E
06H432;
Bichinger L., Pachebat J.A., Gloeckner G., Rajandream M.-A., Sucgang R., Berriman M., Song J., Olsen R., Szafranski K., Xu Q Tunggal B., Kummerfeld S., Madera M., Konfortov B.A., Rivero F. Bankier A.T., Lehmann R., Hamlin N., Davies R., Gaudet P., Fey Pilcher K., Chen G., Saunders D., Sodergren E., Davis P., Ferbornou A., Nie X., Hall N., Anjard C., Hemphill L., Bason N. Farbrother P., Desany B., Just E., Morio T., Rost R., Churcher Cooper J., Haydock S., van Driessche N., Cronin A., Goodhead I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sasaki T., Matsumoto T., Katayose "Oryza sativa nipponbare(GA3) gencolone:P0651G05.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oryza sativa (japonica cultivar-group).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;

Ehrhartoideae; Oryzeae; Oryza.
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05-JUL-2004 (TrEMBLrel. 27, Last sequence up
05-JUL-2004 (TrEMBLrel. 27, Last annotation
                                                                                                                                                                                                                                                                                     Dictyostelium discoideum (Slime mold).
Eukaryota; Mycetozoa; Dictyosteliida; Dictyostelium
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Pfam; PF03962; Mnd1; 1.
SEQUENCE 207 AA; 23900
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InterPro; IPR00564
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (JUN-2003) to the EMBL/GenBank/DDBJ EMBL; AP006528; BAD26517.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Putative MND1 domain containing Name=P0651G05.20;
                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                           NCBI_TaxID=44689;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     40.8%; Score 427; DB 2; Leny... 42.5%; Pred. No. 9.6e-19; A4. Mismatches 71; Indels
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                                                                                                                                                                                                                                                                                                                                                                              Created)
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t P., Fey P.,
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                                                                                                                                Query Match
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Loulseged H., Mungall K., Oliver K., Price C., Quail M.A.,
Urushihara H., Hernandez J., Rabbinowitsch E., Steffen D., Sanders M.,
Ma J., Kohara Y., Sharp S., Simmonds M., Spiegler S., Tivey A.,
Sugano S., White B., Walker D., Woodward J., Winckler T., Tanaka Y.,
Shaulsky G., Schleicher M., Weinstock G., Rosenthal A., Cox E.C.,
Chisholm R.L., Gibbs R., Loomis W.F., Platzer M., Kay R.R.,
Williams J., Dear P.H., Noegel A.A., Barrell B., Kuspa A.,
"The genome of the social amoeba Dictyostelium discoideum.";
Nature 0:0-0(2005).
                                                                                                                                                                                                                                                                                                                                                                                                Submitted (MAR-2005) to the EMBL/GenBank/DDBJ databases.
EMBL; AV811553; AAX27442.1; -; mRNA.
InterPro; IPR0100074; ApoA1_A4_E.
InterPro; IPR010990; Cyt c_b562.
InterPro; IPR011029; DEATH_like.
InterPro; IPR011029; DEATH_like.
InterPro; IPR011991; MINT_LH2.
InterPro; IPR011991; PLAT_LH2.
InterPro; IPR011991; MINT_LH2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hypothetical protein.
Schistosoma japonicum (Blood fluke).
Eukaryota; Metazoa; Platyhelminthes; Trematoda; D
Schistosomatoidea; Schistosomatidae; Schistosoma.
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SCHJA
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i, AAFI01000260; EAL61571.1; -; Genomic_DNA.
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MSVKEVLQSLVDDGMVDCERIGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQ 104
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161 AA; 1
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Pred. No. 7.3e-15;
18; Mismatches 59;
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RA Hollowski G.V., Ussery D., Barrell B.G., Nurse P., Leiner R., Fried for meiotic recombination.
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Q09739;
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"MCp7, a meiosis-specific coiled-coil protein of fission yeast,
associates with Meul3 and is required for meiotic recombination.";
Nucleic Acids Res. 32:3325-3339(2004).
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01-NOV-1995 (Rel. 32, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Meiotic coiled-coil protein 7.
Name=mcp7; ORFNames=SPAC13A11.03;
Schizosaccharomyces pombe (Fission yeast).
Eukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
                                                                                                                                                                                    This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE, FUNCTION, AND INTERACTION WITH MEU13.
PubMed=15210864; DOI=10.1093/nar/9kh654;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN-972
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                                                                                                                                                                                                                                                                                                                                     FUNCTION: Required for meiotic recombination. SUBUNIT: Interacts with meul3.
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   AB189985; BAD42847.1; -; Genomic_
Z54096; CAA90804.1; -; Genomic_Di
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Katinka M.D., Duprat S., Cornillot E., Metenier G., Thomarat F.
Prensier G., Barbe V., Peyretaillade E., Brottier P., Wincker F.
Delbac F., El Alaoui H., Peyret P., Saurin W., Gouy M.,
Weissenbach J., Vivares C.P.;
"Genome sequence and gene compaction of the eukaryote parasite
Encephalitozoon cuniculi.";
Nature 414:450-453(2001).
EMBL; AL590449; CAD258811; -; Genomic DNA.
InterPro; IPR005647; Mnd1.
Pfam; PF03962; Mnd1; 1.
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GeneDB Spombe; SPACIJA11.03; -..
GO; GO:0000785; C:chromatin; IDA.
GO; GO:00007131; P:meiotic recombination; IGI.
GO; GO:0007129; P:synapsis; IMP.
InterPro; IPR005647; Mnd1.
Pfam; PF03962; Mnd1; 1.
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01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Hypothetical protein ECU10 1600.
OrderedLocusNames=ECU10 1600;
Encephalitozoon cuniculi.
                                                                                                                                                                                                                                                                                                                                                               Complete proteome; Hypothetical protein. SEQUENCE 203 AA; 23743 MW; DFE33A65A1A28A42 CRC64;
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QBSUA9_ENCCU PRELIMINARY;
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COILED 77 148 Potential.
SEQUENCE 210 AA; 24224 MW; F4A546F070A37665 CRC64;
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                                                           TSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGRCETEERTRLAKE 126
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   TSNLYWSFASEGIQKKKLRCKELMEECERMSQDICRKREYIENEKMSKHYTEERNELENK 124
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Search completed: February 2, 2006, 12:51:01 Job time : 122 secs

Run

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Minimum DB seq Maximum DB seq

Database

Scoring table: sequence: Title: Perfect score:

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41. Appl
3167, Ap
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1: /cgn2_6/ptodata/1/iaa/5_COMB.pep:*

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3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*

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US-09-621-976-4959
US-09-799-250B-5
US-09-799-250B-721
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APPLICANT: Pia M. Challita-Eid
APPLICANT: Rene S. Hubert
APPLICANT: Steve Chappell Mitchell
APPLICANT: Steve Chappell Mitchell
APPLICANT: Arthur B. Raitano
APPLICANT: Mary Faris
APPLICANT: Daniel E.H. Afar
APPLICANT: Aya Jakobovits
TITLE OF INVENTION: 121PIP1: A TISSUE SPECIFIC PROTEIN
TITLE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS CANCERS
FILE REFERENCE: 129.34US01 (511522003400)
CURRENT APPLICATION NUMBER: US/09/799,250B
CURRENT FILING DATE: 2003-07-14
NUMBER OF SEQ ID NOS: 721
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 205
TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                           KRKFGFEENKIDRTFGIPEDFDYID 205
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US-10-104-047-3167
US-09-103-664A-2
US-08-973-462-8
US-08-272-255-16
PCT-US95-08565-16
US-10-164-955-34
US-09-914-259-65
US-09-914-259-64
US-09-914-259-69
US-09-914-259-69
US-09-914-259-69
US-09-914-259-67
US-09-914-259-47
US-09-914-259-47
US-09-914-259-47
US-09-914-259-57
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Result

Score

Query Match

1047 1011

Length Indels

205; <u>.</u>

Gaps

120 60

120

Pred. No. score grea and is der

948 593.5 349 349 191.5 117 117 111.5 112.5 112.5 112.5 112.5 112.5 112.5 112.5 112.5 112.5 113.5 110.5

11.2 11.2 110.9 110.8 110.7 110.7 110.5 11

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GENERAL INFORMATION:

APPLICANT: Pia M. Challita-Eid

APPLICANT: Steve Chappell Mitchell

APPLICANT: Steve Chappell Mitchell

APPLICANT: Arthur B. Raitano

APPLICANT: Arthur B. Raitano

APPLICANT: Daniel E.H. Afar

APPLICANT: Daniel E.H. Afar

APPLICANT: Aya Jakobovits

TITLE OF INVENTION: 121PIFI: A TISSUE SPECIFIC PROTEIN

TITLE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS CANCERS

FILE REFERENCE: 129.34US01 (511582003400)

CURRENT APPLICATION NUMBER: US/09/799,250B

CURRENT APPLICATION NUMBER: US/09/799,250B

CURRENT FILING DATE: 2003-07-14

INUMBER OF SEO ID NOS: 721

SOFTWARE: FastSQ for Windows Version 4.0

SEQ ID NO 720

LENGTH: 198

TYPE: PRT

ORGANISM: Schizosaccharomyces pombe

US-09-799-250B-720
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US-09-799-250B-4
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                                                                       Query Match 90.5%; Score 948; DB 2; Length 205, Best Local Similarity 89.3%; Pred. No. 1.2e-84; Matches 183; Conservative 10; Mismatches 12; Indels
                                                                                                                                                                                   SEQ ID NO 4
LENGTH: 205
TYPE: PRT
ORGANISM: Mus musculus
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                                                                                                                                                                                                                                                                         APPLICANT: Mary Faris
APPLICANT: Daniel E.H. Afar
APPLICANT: Daniel E.H. Afar
APPLICANT: Aya Jakobovits
TITLE OF INVENTION: 121P1F1: A TISSUE SPECIFIC PROTEIN
TITLE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS CANCER
FILE REFERENCE: 129.34US01 (511582003400)
CURRENT APPLICATION NUMBER: US/09/799,250B
CURRENT FILING DATE: 2003-07-14
NUMBER OF SEQ ID NOS: 721
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Pia M. Challita-Eid
APPLICANT: Rene S. Hubert
APPLICANT: Steve Chappell Mito
APPLICANT: Arthur B. Raitano
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Steve Chappell Mitchell Arthur B. Raitano
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RESULT 5
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                                                                                                                                                                      APPLICANT: Pia M. Challita-E
APPLICANT: Rene S. Hubert
APPLICANT: Steve Chappell M
APPLICANT: Arthur B. Raitan
APPLICANT: Mary Faris
APPLICANT: Mary Faris
APPLICANT: Aya Jakobovite
TITLE OF INVENTION: 12191F1: A TISSUE SPECIFIC PROTEIN TITLE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS CANCER FILE REFERENCE: 129.34US01 (511582003400) CURRENT APPLICATION NUMBER: US(99/799,250B CURRENT FILING DATE: 2003-07-14 NUMBER OF SEQ ID NOS: 721 SOFTWARE: PRACESQ for Windows Version 4.0 SEQ ID NO 5 LENGTH: 200
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Patent No. 6639063
GENERAL INFORMATION:
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SOFTWARE: Patent.pm
SEQ ID NO 4959
LENGTH: 127
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                                                                                                                                                                                                                                                                                                                            Sequence 5, Application US/09799250B Patent No. 6924358
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Best Local Similarity
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APPLICANT: Jobert, S.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: ESTs and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                               ----IKLSGMQEER 127
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                                                                                                                                                                                                                               Rene S. Hubert
Steve Chappell Mitchell
Arthur B. Raitano
                                                                                                                                                                                            Mary Faris
Daniel E.H. Afar
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Pred. No. 2.4e-50;
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RESULT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: 121P1F1: A TISSUE SPECIFIC PROTEIN
TITLE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS CANCERS
FILE REFERENCE: 129.34US01 (51158203400)
CURRENT APPLICATION NUMBER: US/09/799,250B
CURRENT FILING DATE: 2003-07-14
NUMBER OF SEQ ID NOS: 721
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 721
                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Pia M. Challita-Eid
APPLICANT: Rene S. Hubert
APPLICANT: Steve Chappell Mito
APPLICANT: Arthur B. Raitano
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT ORGANISM: Schizosaccharomyces pombe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Schizosaccharomyces pombe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 200
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                                                                                                                                                                                                                                                                                                                                                                                     Match 33.3%; Score 349; DB 2; Length 200; Local Similarity 41.6%; Pred. No. 3.2e-26; es 84; Conservative 36; Mismatches 76; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         122 RLAKELSSLRD-QREQLKAEVEKYKDCDPQVVEBIRQANKVAKEAANRWTDNIFAIKSWA 180
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                                                                                                               181 KRKFGFEENKIDRTFGIPEDFD 202
                                                                                                                                                                                         122 RLAKELSSLRD-QREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWA 180
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. 6924358
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                                                                                                                                                                                                                                                                                                           1 KGLSLAEKRRLEAIFHDSKDFFQLKEVEKLGSK-KQIVLQTVKDVLQSLVDDNIVKTEK
                                                                                                                                                                                                                                                                                                                                   5 KGLSAEEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMVDCER 64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KGLSAEEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMVDCER 64
                                                                                                                                                     QYTLELLHAKESELKLLKTQLSNLNHCNPETFELKNENTKKYMEAANLWTDQIHTLIAFC
                                                                                                                                                                                                                                                                     IGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQKSI--EKAKIGRCETE-ERT 121
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Steve Chappell Mitchell
Arthur B. Raitano
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                                                                                                                                                                                                                                                                                                                                                                                       6,
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                                                                                                                                                                                                                                 119
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Sequence 14485, Application US/09248796A

Patent No. 6747137

GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
ITITLE OF INVENTION: NUCLEIC ACID AND ACID SEQUENCES RELATING TO C
ITITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132

CURRENT APPLICATION NUMBER: US/09/248,796A

CURRENT FILING DATE: 1999-02-12

PRIOR APPLICATION NUMBER: US 60/074,725

PRIOR PILING DATE: 1998-02-13

PRIOR APPLICATION NUMBER: US 60/096,409

PRIOR PILING DATE: 1998-08-13

PRIOR APPLICATION NUMBER: US 60/096,409

PRIOR PILING DATE: 1998-08-13
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                                                                                                                                    US-09-538-092-1321
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                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 1387
SOFTWARE: CuraPatSeqFormatter Version
SEQ ID NO 1321
LENGTH: 1270
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SEQ ID NO 14485
LENGTH: 260
                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1321, Application US/09538092 Patent No. 6753314
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Best Local 9
                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 15966-542
CURRENT APPLICATION NUMBER: US/09/538,092
CURRENT FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: 60/127,352
PRIOR FILING DATE: 1999-04-01
PRIOR APPLICATION NUMBER: 60/178,965
PRIOR FILING DATE: 2000-02-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Giot, Loic
APPLICANT: Mansfield, Traci A.
TITLE OF INVENTION: Protein-Protein Complexes
                                                                                                                                                   PEATURE: misc_feature
NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: Polypeptide Accession Number Q14203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Candida albicans
                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18.3%; Score 191.5; DB 2
Local Similarity 27.1%; Pred. No. 1.1e-10;
1e8 59; Conservative 47; Mismarcher
                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               167 NRWTDNIFAIKSWAKRKFGFEENKIDRTFGIPEDFDYI 204
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       115 C--ETEERTRIAKEISSIR-----DQREQIKAEVEKYKDCDPQVVEBIRQANKVAKEAA 166
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2 SKKKGLSAEEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMVD
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                                             Conservative
                                                                  12.8%; Score 134.5; DB 2 23.4%; Pred. No. 0.00032;
                                             43;
                                             Mismatches 106;
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                                                                                     DB 2;
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                                                                                       Length 1270;
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Patent No. 6232460
                                                                                                                                                                             Matches
                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                            TELEFAX: (212) 752-5958
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                        NAME: Hanson, No. 6232460man D. REGISTRATION NUMBER: 30,946
REFERENCE/DOCKET NUMBER: LUD 54
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/892,702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Wordperfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 15-July-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                  LENGTH: 976 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/104,324B FILING DATE: 25-June-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE:
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                                                                                                                                                                                             ocal Similarity
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                                                                                                                   413 QKKSSELEEMTKLTNNKEVELEBLKKVLGEKETLLYBNKQFEKIAEELKG-TEQELIGLL
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                                                                                                                                                3 KKKGLSAEE--KRTRMMEI-FSETKDV-----FOLKDLEKIAPKEKGITAMSVKEVL
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                                                          QAREKEVHDLEIQLTAITTSEQYYSKEVKDLKTELENEKLKNTELTSHCNKLSLBNKELT
                                                                                       QSL---VDDGMVDCERIGTSNYYWAFPSKAL-----
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QETSDMTLELKNQQEDINNNKKQEERMLKQIE--NLQETETQLRNELBYVREELKQKRDE
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666 Fifth Avenue
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Methods For Diagnosis And Treating Cancers,
And Methods For Identifying Pathogenic Markers In
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                                                                                                                                                                                          11.2%; Score 117; DB 2 23.9%; Pred. No. 0.012;
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                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                       DB 2; Length 976;
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                                                                                      -----HARKHKLEVLE--
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                                                                                                                                                                             Gaps
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                                                                                                                                  US-09-914-259-55
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US-09-538-092-1339
; Sequence 1339, Applice
; Patent No. 6753314
; Patent No. 6753314
; GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/09/538,092

CURRENT FILING DATE: 2000-03-29

PRIOR APPLICATION NUMBER: 60/127,352

PRIOR FILING DATE: 1999-04-01

PRIOR APPLICATION NUMBER: 60/178,965

PRIOR FILING DATE: 2000-02-01

NUMBER OF SEQ ID NOS: 1387

SOFTWARE: CuraPatSeqFormatter Version 0.9

SEQ ID NO 1339
                                                                                                                                   Sequence 55, Application Patent No. 6495336 GENERAL INFORMATION:
               APPLICANT: Makowski, Lee
APPLICANT: Hyman, Paul
APPLICANT: Williams, Mark
TITLE OF INVENTION: STAGED ASSEMBLY OF NANOSTRUCTURES
FILE REFERENCE: 8471-010-999
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CURRENT APPLICATION NUMBER: US/09/914,259
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ORGANISM: Homo sapiens
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NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: Polypeptide Accession Number Q15431
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                                                                                                                                                                                                                                                                                                                  650 LESAKQKFGEITDTYQKEIEDKKI 673
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CURRENT APPLICATION NUMBER: US/09/914,259
; CURRENT FILING DATE: 2000-11-21
; NUMBER OF SEQ ID NOS: 180
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 62
; SEQ ID NO 62
; LENGTH: 284
; TYPE: PRT
; ORGANISM: Rattus norvegicus
US-09-914-259-62
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NUMBER OF SEQ ID NOS: 1

SOFTWARE: FASTSEQ for W

SEQ ID NO 55

LENGTH: 284

TYPE: PRT

ORGANISM: Sus scrofa
US-09-914-259-55
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US-09-914-259-62
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RESULT 13
US-09-248-796A-20275
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APPLICANT: Makowski, Lee
APPLICANT: Hyman, Paul
APPLICANT: Williams, Mark
TITLE OF INVENTION: STAGED ASSEMBLY OF NANOSTRUCTURES
FILE REFERENCE: 8471-010-999
                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                   161 KYEEVARKLVIIESDLERAEERAELSEGKC-----AELEEELKTVTNNLKSLEAQAEKYS
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                                                                         216 QKEDKYEEEIKVLSDKLKEAETR--
                                                                                                                                                                                                                              104 ---ERLATALOKLEBAEKAADESERGMKVIESRAQKDEEKMEIQEIQLKEAKHIAEDADR 160
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                                                                                                                                                                                                                                                                                                                                          2 SKKKGLSA-EEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVK-EVLQSLVDDGM 59
                                                                                                                                                                                                                                                                VDCERIGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQK--------
                                                                                                             DCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWAKRKFGFEENKID 192
                                                                                                                                                                                                                                                                                                    AKEKLLRASEDERDRVLEELHKAEDSLLAAD-ETAAKAEADVASLNRRIQLVEEELDRAQ 103
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180
Windows Version
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21.1%; pred. No. 0.0048;
40. Mismatches 78;
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Pred. No. 0.0043;
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                                                                           ----AEFAERSVTKLEKSID 254
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FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
INUMBER OF SEQ ID NOS: 28208
SEQ ID NO 20275
LENGTH: 630
TYPE: PRT
ORGANISM: Candida albicans
US-09-248-796A-20275
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                                                                                                                                                                                                                                 ; LENGTH: 284
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-914-259-43
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US-09-914-259-43
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CURRENT FILING DATE: 2000-11-21
NUMBER OF SEQ ID NOS: 180
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 43
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 43, Application US/09914259
Patent No. 6495336
GENERAL INFORMATION:
APPLICANT: Makowski, Lee
APPLICANT: Hyman, Paul
APPLICANT: Williams, Mark
TITLE OF INVENTION: STAGED ASSEMBLY OF NANOSTRUCTURES
FILE REFERENCE: 8471-010-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20275, Application US/09248796A Patent No. 6747137
GENERAL INFORMATION:
                                                                                                                                                      Matches
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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133 R----ALKDEEKMELQEI--
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                                                                                                             1 MSKKKGLSAEEK---RTRMMEIFSETKDVFQ-----LKDLEKIAPK-EKGITAMSV 47
                                                                                                                                                    th 10.7%; Score 112; DB 2; l Similarity 23.5%; Pred. No. 0.0068; 52; Conservative 41; Mismatches 72;
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                                                                        N---DN-----SGAKKELLEKVSKLE 344
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -EKAKIGRCETEERTRLAKELSSLRDQREQLKAEVEKYKDCDPQVVBEIRQANKVAKEAA 166
                                   KEVLQSLVDDGMVDCERIGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQKSI 107
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---QLKEAKHIAEEADRKYEEVARKLVIIEGDL 176
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GENERAL INFORMATION: Glot, Loic
APPLICANT: Glot, Loic
APPLICANT: Mansfield, Traci A.

TITLE OF INVENTION: Protein-Protein Complexes and Method of Using Same
FILE REPERENCE: 15966-542
CURRENT PAPLICATION NUMBER: US/09/538,092
CURRENT FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: 60/173,352
PRIOR FILING DATE: 1999-04-01
PRIOR FILING DATE: 2000-02-01
NUMBER OF SEQ ID NOS: 1387
SOFTWARE: CuraPatSeqFormatter Version 0.9
SEQ ID NO 918
LENGTH: 1937
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
NAME/KEX: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: Polypeptide Accession Number P13535
US-09-538-092-918
Search completed: February
Job time : 48 secs
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US-09-538-092-918
; Sequence 918, Application US/09538092
; Patent No. 6753314
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                                                                                              1536 VEQEK-CEIQAALEEAEASLEHEEGKILRIQLELNQVKSEVDRKIAEKDEEIDQ 1588
                                                                                                                                                                                                  1476 ESRSISTELFKVKNVYEESLDQLETLRRENKNIQQEISDLTEQIAEGGKQIHELEKIKKQ 1535
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                                                                                                                                                                                                                                                                                                                                                                                                       1361 QRALSKANSEVAQWRTKYETDAIQRTEELEEAKKKLAQRLQEAEEHVEÄVNAKCASLEKT 1420
                                                                                                                                                141 VEKYKDCDPQVVEEIRQANKVAKEA-ANRWTDNIFAIKSWAKRKFGFEENKIDR 193
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                                                                                                                                                                                                                                                    99 KHASLQKSIEKAK-----IGRCET--EERTRLAKELSSLRDQ-----REQLKAE 140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 KKGLSABEKRTRMMBIFSETKDVFQLKDLBKIAPK------BKGITAMS------V 47
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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US-11-087-190-61
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US-11-125-085-2
US-10-087-190-44
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Sequence 3, Appli
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Sequence 21, Appl
Sequence 22, Appl
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Sequence 67, Appl
Sequence 68, Appl
Sequence 69, Appl
Sequence 69, Appl
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Sequence 3, Application US/10087190 Publication No. US20030223997A1
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287	329.5	349	349	349	349	394	427	434	465	465	465	584	584	584	591	591	591
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US-10-106-698-5443	US-10-450-763-42700	US-11-125-805-721	US-11-125-805-5	US-10-087-190-25	US-09-799-250-5	US-10-437-963-162901	US-10-437-963-162899	US-10-425-115-253013	US-10-087-190-46	US-10-087-190-15	US-10-087-190-5	US-10-087-190-47	US-10-087-190-16	US-10-087-190-7	US-10-087-190-55	US-10-087-190-54	US-10-087-190-17
Sequence 5443, Ap	Sequence 42700, A	Sequence 721, App	Sequence 5, Appli	Sequence 25, Appl	Sequence 5, Appli	Sequence 162901,	Sequence 162899,	Sequence 253013,	Sequence 46, Appl	Sequence 15, Appl	Sequence 5, Appli	Sequence 47, Appl	Sequence 16, Appl	Sequence 7, Appli	Sequence 55, Appl	Sequence 54, Appl	Sequence 17, Appl

ALIGNMENTS

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TITLE OF INVENTION: 12191F1: A TISSUE SPECIFIC PROTEIN
TITLE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS CANCERS
FILE REFERENCE: 129.34US01
CURRENT APPLICATION NUMBER: US/09/799,250
CURRENT FILING DATE: 2001-02-08
NUMBER OF SEQ ID NOS: 719
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 205
TYPE: PRT
ORGANISM: Homo sapiens
US-09-799-250-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 2, Application US/09799250
Publication No. US20030032087A1
GENERAL INFORMATION:
APPLICANT: Pia M. Challita-Eid
APPLICANT: Rene S. Hubert
APPLICANT: Steve Chappell Mitchell
APPLICANT: Arthur B. Raitano
APPLICANT: Mary Faris
APPLICANT: Daniel B.H. Afar
APPLICANT: Aya Jakobovits
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Best Local Similarity 100.0%;
Matches 205; Conservative 0
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KRKFGFEENKIDRTFGIPEDFDYID
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Pred. No. 4e-78;
0; Mismatches
205
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APPLICANT: Faris, Mary
APPLICANT: Afar, Daniel E. H.
APPLICANT: Ge, Wangmao
APPLICANT: Jakobovitz, Aya
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: ENTITLED 121P1F1 USEFUL IN TREATMENT AND
FILE REFERENCE: 51158-20034.20
CURRENT APPLICATION NUMBER: US/10/087,190
CURRENT FILING DATE: 2003-01-28
PRIOR APPLICATION NUMBER: US 09/779,250
PRIOR APPLICATION NUMBER: US 09/779,250
PRIOR APPLICATION NUMBER: US 09/779,250
PRIOR FILING DATE: 2001-03-05
VUMBER OF SEQ ID NOS: 69
US-10-087-190-14
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LENGTH: 205
TYPE: PRT
ORGANISM: Homo Sapiens
                                                                                         SEQ ID NO 14
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Agensys, Inc.
APPLICANT: Challita-Eid, Pia M.
APPLICANT: Hubert, Rene S.
APPLICANT: Raitano, Arthur B.
APPLICANT: Faris, Mary
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TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: ENTILED 121P1F1 USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: S1158-20034.20
CURRENT APPLICATION NUMBER: US/10/087,190
CURRENT FILING DATE: 2003-01-28
PRIOR APPLICATION NUMBER: US 09/779,250
PRIOR PILING DATE: 2001-03-05
NUMBER OF SEQ ID NOS: 69
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APPLICANT: Challita-Eid, Pia M.
APPLICANT: Hubert, Rene S.
APPLICANT: Raitano, Arthur B.
                                                                                                              NUMBER OF SEQ ID NOS: 69
SOFTWARE: FastSEQ for Windows Version 4.0
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                   LENGTH: 205
TYPE: PRT
ORGANISM: Homo Sapiens
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Afar, Daniel E. H.
Ge, Wangmao
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US-10-087-190-20
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                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 20
LENGTH: 205
TYPE: PRT
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APPLICANT: Challita-Eid, Pis
APPLICANT: Hubert, Rene S.
APPLICANT: Raitano, Arthur E
APPLICANT: Faris, Mary
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Best Local Similarity
Matches 205; Conserv
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Matches 205;
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APPLICANT: Jakobovitz, Aya
APPLICANT: Jakobovitz, ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: ENTITLED 121P1F1 USEFUL IN TREATMENT AND DETECTION OF
FILE REFERENCE: 51158-20034-20
CURRENT EPPLICATION NUMBER: US/10/087,190
CURRENT FILING DATE: 2003-01-28
FRIOR APPLICATION NUMBER: US 09/779,250
PRIOR PILING DATE: 2001-03-05
NUMBER OF SEQ ID NOS: 69
SOFTWARE: FastSEQ for Windows Version 4.0
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                          KRKFGFEENKIDRTFGIPEDFDYID 205
                                                                                                      TRLAKELSSIRDQREQLKAEVEKYKDCDPQVVBEIRQANKVAKEAANRWTDNIFAIKSWA 180
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  KRKFGFEENKIDRTFGIPEDFDYID
                                                                            TRLAKELSSLRDQREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWA
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Afar, Daniel E. H.
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                                                                                                                                                                                                                                                                                                               100.0%; Score 1047; DB 4;
100.0%; Pred. No. 4e-78;
ative 0; Mismatches 0;
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US-10-087-190-21 ; Sequence 21, Application US/10087190

RESULT 5

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APPLICANT: Afar, Daniel E. H.
APPLICANT: Ge, Wangmao
APPLICANT: Jakobovitz, Aya
APPLICANT: Jakobovitz, Aya
ITILE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: ENTITLED 121P1F1 USEFUL IN TREATMENT AND
FILE REFERENCE: 51158-20034.20
CURRENT APPLICATION NUMBER: US/10/087,190
CURRENT FILING DATE: 2003-01-28
PRIOR APPLICATION NUMBER: US 9/779,250
PRIOR APPLICATION NUMBER: US 9/779,250
PRIOR FILING DATE: 2001-03-05
NUMBER OF SEQ ID NOS: 69
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 22
SEQ ID NO 22
                                                                                                                                                                                                                                                                                                                                                                            Sequence 22, Application US/10087190
Publication No. US20030223997A1
GENERAL INFORMATION:
APPLICANT: Agensys, Inc.
APPLICANT: Challita-Eid, Pia M.
APPLICANT: Hubert, Rene S.
APPLICANT: Raitano, Arthur B.
APPLICANT: Faris, Mary
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US-10-087-190-22
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ORGANISM: Homo Sapiens US-10-087-190-22
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US-10-087-190-21
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LENGTH: 205
TYPE: PRT
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APPLICANT: Ge, Wangmao
APPLICANT: Jakobovitz, Aya
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: ENTITLED 121P1F1 USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20034.20
CURRENT APPLICATION NUMBER: US/10/087,190
CURRENT APPLICATION NUMBER: US/09/779,250
PRIOR APPLICATION NUMBER: US/09/779,250
PRIOR APPLICATION NUMBER: US/09/779,250
NUMBER OF SEQ ID NOS: 69
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APPLICANT: Challita-Eid, Pia M.
APPLICANT: Hubert, Rene S.
APPLICANT: Raitano, Arthur B.
APPLICANT: Faris, Mary
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                                              LENGTH: 205
TYPE: PRT
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Local Similarity 100.0%;
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Afar, Daniel E. H.
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Afar, Daniel E. H.
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US-10-087-190-61
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Best Local Similarity
Matches 205; Conserv
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                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 09/779,250
PRIOR FILING DATE: 2001-03-05
NUMBER OF SEQ ID NOS: 69
SOFTWARE: FRANCES
                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Jakobovitz, Aya
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: ENTITLED 121P1F1 USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20034.20
CURRENT APPLICATION NUMBER: US/10/087,190
CURRENT FILING DATE: 2003-01-28
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APPLICANT: Challita-Eic
APPLICANT: Hubert, Rene
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                                                                                                                                                                                                                                                                                                                                                    LENGTH: 205
TYPE: PRT
ORGANISM: Homo Sapiens
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KRKFGFEENKIDRTFGIPEDFDYID 205
                   KRKFGFEENKIDRTFGIPEDFDYID 205
                                                                TRLAKELSSLRDQREQLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWA
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Afar, Daniel E. H.
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RESULT 8 US-10-408-765A-1821

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APPLICANT: Ghosh, Soumitra S.
APPLICANT: Fahy, Eoin D.
APPLICANT: Fahy, Eoin D.
APPLICANT: Gibson, Bradford W.
APPLICANT: Gibson, Bradford W.
APPLICANT: Taylor, Steven W.
APPLICANT: Warnock, Dale E.
TITLE OF INVENTION: TARGETS FOR THERAPEUTIC INTERVENTION
TITLE OF INVENTION: IDENTIFIED IN THE MITOCHONDRIAL PROTEOME
FILE REFERENCE: 660088.465
CURRENT APPLICATION NUMBER: US/10/408,765A
CURRENT FILING DATE: 2003-04-04
NUMBER OF SEQ ID NOS: 3077
SECTIMARE: FastSEQ for Windows Version 4.0
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US-11-125-805-2
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; LENGTH: 205
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-408-765A-1821
                                                                US-11-125-805-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/11125805 Publication No. US20050208059A1 GENERAL INFORMATION:
                                                                                 SEQ ID NO 2
LENGTH: 205
TYPE: PRT
ORGANISM: Homo sapiens
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Publication No. US20
GENERAL INFORMATION:
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Query Match
Best Local Similarity
                                                                                                                                                                                APPLICANT: AVA JAKODOVILE

TITLE OF INVENTION: 121P1F1: A TISSUE SPECIFIC PROTEIN

TITLE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS CANCERS

FILE REFERENCE: 129.34U801 (511.582003400)

CURRENT APPLICATION NUMBER: US/11/125,805

CURRENT FILING DATE: 2005-05-09

PRIOR APPLICATION NUMBER: US/09/799,250B

PRIOR APPLICATION NUMBER: US/09/799,250B

PRIOR FILING DATE: 2003-07-14

NUMBER OF SEQ ID NOS: 721
                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Pia M. Challita-Eid
APPLICANT: Rene S. Hubert
APPLICANT: Steve Chappell Mitchell
APPLICANT: Arthur B. Raitano
                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0
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Daniel E.H. Afar
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100.0%;
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Score 1047; DB Pred. No. 4e-78;
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                      Length 205;
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APPLICANT: Afar, Daniel E. H.
APPLICANT: Ge, Wangmao
APPLICANT: Jakobovitz, Aya
ITITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
ITITLE OF INVENTION: ENTITLED 121P1F1 USEFUL IN TREATMENT AND
FILE REFERENCE: 51158-20034.20
CURRENT APPLICATION NUMBER: US/10/087,190
CURRENT FILING DATE: 2003-01-28
PRIOR APPLICATION NUMBER: US 09/779,250
PRIOR FILING DATE: 2001-03-05
NUMBER OF SEQ ID NOS: 69
COPTMEND: TSICED FOR MINISTREED TO MAINTAIN TO THE PRIOR APPLICATION NUMBER: US 09/779,250
                                                                                                                                                                                                                                                                                                                                                          ; LENGTH: 206
; TYPE: PRT
; ORGANISM: Homo 9
US-10-087-190-44
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Best Local Similarity
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APPLICANT: Challita-Eid, P
APPLICANT: Hubert, Rene S.
APPLICANT: Raitano, Arthur
APPLICANT: Faris, Mary
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                                                                                                                                                                                                                 MSKKKGLSABEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMV
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                  AKRKFGFEENKIDRTFGIPEDFDYID 205
                                                                                                                                           DCERIGISNYYWAFPSKALHARKHKLEVLESQQLSEGSQKHASLQKSIEKAKIGRCETEE
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                                                                    RTRLAKELSSLRDQREQLKAEVEKYKDCDPQVVBEIRQANKVAKEAANRWTDNIFAIKSW
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Pred. No. 3e-77;
0; Mismatches
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RESULT 11
US-10-087-190-45
; Sequence 45, Application US/10087190
; Publication No. US20030223997A1
; GENERAL INFORMATION:

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APPLICANT: Afar, Daniel E. H.

APPLICANT: Ge, Wangmao
APPLICANT: Jakobovitz, Aya
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: SWTITLED 121P1F1 USEFUL IN TREATMENT AND DETECTION OF CANCE
FILE REFERENCE: 51158-20034.20
CURRENT FILLORISM: USO3-01-28
CURRENT FILLORISM: USO3-01-28
PRIOR APPLICATION NUMBER: US 09/779,250
PRIOR FILLORISM: LONG: 69
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 45
LENGTH: 206
TYPE: PRT
ORGANISM: Homo Sapiens
US-10-087-190-45

OG OR: SCOTE 1036.5; DB 4; Length 206;
                                                                                         APPLICANT: Agensys, Inc.
APPLICANT: Agensys, Inc.
APPLICANT: Hubbert, Rene S.
APPLICANT: Hubbert, Rene S.
APPLICANT: Raitano, Arthur B.
APPLICANT: Raitano, Arthur B.
APPLICANT: Afar, Daniel E. H.
APPLICANT: Ge, Wangmao
APPLICANT: Ge, Wangmao
APPLICANT: Jakobovitz, Aya
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
CURRENT APPLICATION NUMBER: US/10/087,190
CURRENT APPLICATION NUMBER: US/10/087,190
CURRENT FILING DATE: 2003-01-28
PRIOR APPLICATION NUMBER: US 09/779,250
PRIOR FILING DATE: 2001-03-05
NUMBER OF SEQ ID NOS: 69
SOFTWARE: FastSEQ for Windows Version 4.0
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US-10-087-190-24
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                                    US-10-087-190-24
                                                  LENGTH: 198
• TYPE: PRT
ORGANISM: Homo Sapiens
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Query Match
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APPLICANT:
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Afar, Daniel E. H.
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   96.6%;
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Pred. No. 3e-77;
0; Mismatches
     Score 1011;
     DB
     4
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   Length 198;
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   RESULT 14
US-10-687-190-13
; Sequence 13, Application US/10087190
; Publication No. US20030223997A1
; GENERAL INFORMATION:
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APPLICANT: Daniel B.H. Afar
APPLICANT: APA JAKOBOVILE
APPLICANT: APA JAKOBOVILE
TITLE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS CANCERS
FILE REFERENCE: 129.34US01 (511582003400)
CURRENT APPLICATION NUMBER: US/11/125,805
CURRENT FILING DATE: 2005-05-09
PRIOR APPLICATION NUMBER: US/9/799,250B
PRIOR APPLICATION NUMBER: US/09/799,250B
PRIOR FILING DATE: 2003-07-14
NUMBER OF SEQ ID NOS: 721
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                   ; SEQ ID NO 720
; LENGTH: 198
; TYPE: PRT
; ORGANISM: Schizosaccharomyces pombe
US-11-125-805-720
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US-11-125-805-720
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Publication No. US20050208059A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Pia M. C
APPLICANT: Rene S:
APPLICANT: Steve C
APPLICANT: Arthur
APPLICANT: Mary Fa
APPLICANT: Danial
                                                                                                                                                                                                                                                                                                                                  Query Match 96.6%; Score 1011; DB 6;
Best Local Similarity 100.0%; Pred. No. 3.5e-75;
Matches 198; Conservative 0; Mismatches 0;
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Steve Chappell Mitchell
Arthur B. Raitano
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GFEENKIDRTFGIPEDFD 198
                     GFEENKIDRTFGIPEDFD 202
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Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                            Length 198;
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APPLICANT: Faits, Mary
APPLICANT: General E. H.
APPLICANT: General E. H.
APPLICANT: General E. H.
APPLICANT: Jakobovitz, Aya
ITILE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: ENTITLED 121P11 USEFUL IN TREATMENT AND DETECTION OF CANC
FILE REPERENCE: 51158-20034.20
CURRENT APPLICATION NUMBER: US/10/087,190
CURRENT FILING DATE: 2003-01-28
PRIOR APPLICATION NUMBER: US 09/779,250
PRIOR APPLICATION NUMBER: US 09/779,250
PRIOR FILING DATE: 2001-03-05
NUMBER OF SEQ ID NOS: 69
SOFTWARE: FRASCECQ for Windows Version 4.0
SEQ ID NO 13
                                                                                                                                   APPLICANT: Agensys, Inc.
APPLICANT: Challita-Eid, Pia M.
APPLICANT: Hubert, Rene S.
APPLICANT: Hubert, Rene S.
APPLICANT: Raitano, Arthur B.
APPLICANT: Faris, Mary
APPLICANT: Afar, Daniel E. H.
APPLICANT: Ge, Wangmao
APPLICANT: Jakobovitz, Aya
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN
TITLE OF INVENTION: ENTITLED 121PLF1 USEFUL IN TREATMENT AND
FILE REFERENCE: 51158-20034.20
CURRENT FILING DATE: 2003-01-28
PRIOR APPLICATION NUMBER: US/10/087,190
CURRENT FILING DATE: 2003-01-28
PRIOR APPLICATION NUMBER: US 09/779,250
PRIOR FILING DATE: 2001-03-05
NUMBER OF SEG ID NOS: 69
SOFTWARE: FastSEQ for Windows Version 4.0
                                          ; LENGTH: 190
; TYPE: PRT
; ORGANISM: Homo S
US-10-087-190-19
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; TYPE: PRT
; ORGANISM: Homo Sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19, Application US/10087190 Publication No. US20030223997A1
Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               181 GIPEDFDYID 190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       76 SKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGRCETEERTRLAKELSSLRDQRE 135
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Challita-Eid, Pia M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            QLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWAKRKFGFEENKIDRTF
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                                                                      Sapiens
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93.1%; Score 975;
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Length 190;
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181 GIPEDFDYID 190
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                                                                                                                      61 SKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGRCETEERTRLAKELSSLRDQRE 120
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                                                                           QLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWAKRKFGFEENKIDRTF 195
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                                                         QLKAEVEKYKDCDPQVVEEIRQANKVAKEAANRWTDNIFAIKSWAKKKFGFEENKIDRTF 180
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Search completed: February 2, 2006, 12:56:14
Job time: 173 secs

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Result
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1: /cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/USO9_NEW_PUB.pep:*
6: /cgn2_6/ptodata/1/pubpaa/USO1_NEW_PUB.pep:*
7: /cgn2_6/ptodata/1/pubpaa/USO1_NEW_PUB.pep:*
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GenCore version 5.1.7
(c) 1993 - 2006 Biocceleration
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US-11-124-368A-214
US-11-124-368A-215
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US-11-055-309A-10
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US-10-821-234-1136
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Sequence 59, Appl
Sequence 3, Appli
Sequence 52, Appl
Sequence 214, App
Sequence 215, Appl
Sequence 54, Appl
Sequence 54, Appl
Sequence 183, App
Sequence 1632, Ap
Sequence 175, App
Sequence 175, App
Sequence 44, Appl
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10, Appl
50, Appl
23, Appl
1171, Ap
1143, Ap
136, App
901, App
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8.0	8.0	8.0	8.1	8.1	8.1	8.1	8.1	8.1	8.1	8.2	8.4	8.4	8.4	8.4	8.4	8.4	8.4	8.4
1652	663	663	3116	2480	2107	1641	448	693	227	1663	1732	5464	5415	5406	5335	3960	3803	1299
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US-10-995-561-663	US-11-196-475-78	US-11-196-475-70	US-10-995-561-826	US-10-995-561-825	US-10-995-561-827	US-10-877-346-40	US-10-793-626-1728	US-10-873-528-185	US-11-215-658-12	US-10-055-877-148	US-10-055-877-147	US-10-995-561-775	US-10-995-561-779	US-10-995-561-774	US-10-995-561-777	US-10-995-561-771	US-10-995-561-773	US-10-821-234-1145
Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence
663, App	78, Appl	70, Appl		•	•		1728, Ap	185, App	12, Appl	148, App	147, App						773, App	

ALIGNMENTS

RESULT 1 US-11-155-288-20 US-11-155-288-20 Sequence 20, Application US/11155288 Publication No. US20060008466A1 Publication No. US20060008466A1 GENERAL INFORMATION: APPLICANT: Chiang, Chih-Sheng APPLICANT: Simard, John J.L. TITLE OF INVENTION: COMBINATIONS OF TUMOR-ASSOCIATED TITLE OF INVENTION: ANTIGENS IN DIAGNOSTICS FOR VARIOUS TYPES OF CANCERS FILE REFERENCE: MANNK.050A CURRENT APPLICATION NUMBER: US/11/155,288 CURRENT APPLICATION NUMBER: 60/580,969 PRIOR APPLICATION NUMBER: 60/580,969 PRIOR FILING DATE: 2004-06-17 NUMBER OF SEC ID NOS: 40 ; LENGTH: 976 ; TYPE: PRT ; ORGANISM: Homo sapiens US-11-155-288-20 Query Match Best Local Similarity 23.9%; SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 20 Matches 472 650 LESAKOKFGEITDTYQKEIEDKKI 532 QETSDMTLELKNQQEDINNNKKQEERMLKQIE--NLQETETQLKNELEYVREELKQKRDE 413 QKKSSELEEMTKLTNNKEVELEELKKVLGEKETLLYENKQFEKTAEELKG-TEQELIGLL 91 52 QSL---VDDGMVDCERIGTSNYYWAFPSKAL-------HARKHKLEVLE--3 KKKGLSAEE--KRTRMMEI-FSETKDV-----FQLKDLEKIAPKEKGITAMSVKEVL 63; Conservative LKAEVEKYKD-CD-------PQVVEEIRQANKVAKE--AANRWTDNIFAIK-------SWAKRKFG-----FEENKI 191 VKCKLDKSEENCHNLRKQVENKNKYIBELQQENKALKKKGTAESKQLNVYEIKVNKLELE 649 -----SQLSEGSQKHASLQKSIEKAKIGRCETEERTRLAKELSSLRDQREQ QAREKEVHDLEIQLTAITTSEQYYSKEVKDLKTELENEKLKNTELTSHCNKLSLENKELT 531 ; Score 117; DB 7; Length 976; pred. No. 0.094; 38; Mismatches 85; Indels 673 78; Gaps 589 136 90 471 51 13

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RESULT 2

Sequence 59, Applica Publication No. US20 GENERAL INFORMATION:

Application US/11019711 TO. US20060009634A1

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                                                                                                                                        ; OTHER INFORMATION: ; OTHER INFORMATION: US-11-019-711-59
                                                                      Matches
                                                                                      Query Match
Best Local Similarity
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APPLICANT:
APPLICANT:
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APPLICANT: Eisen, Andrew J
TITLE OF INVENTION: Proteins and Nucleic Acids Encoding
FILE REFERENCE: 21402-235
                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 2001-09-12
NUMBER OF SEQ ID NOS: 227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/11/019,711
CURRENT FILING DATE: 2004-12-21
PRIOR APPLICATION NUMBER: US/10/037,417
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                                                                                                                                                                                                                                          ENGTH: 860
                                                                                                                                                                                                                                                              ID NO 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 60/260,018
FILING DATE: 2001-01-05
APPLICATION NUMBER: 60/260,360
FILING DATE: 2001-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/272,411
FILING DATE: 2001-02-28
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                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 60/318,700
                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/318,405 FILING DATE: 2001-09-10
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 60/305,060
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/272,817
                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 2001-07-
                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 2001-07-05
                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/303,231
                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 2001-05-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/291,186
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                                     13
 RKKHQDAINELSE--QIEQLQKQKAK----AEKEKSQLQAEVDDLLAQLDSI-----
                                 RTRMMBIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMVDCERIGTSNYYW
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Padigaru, Muralidhara
Taupier Jr, Raymond J
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Boldog, Ferenc L
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Grosse, William M
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Tchernev, Velizar T
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Sciore, Paul
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Vernet, Corine A.M.
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                                                                                                                                                      Description of Artificial Sequence: Mysosin Tail Consensus Sequence
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                                                                                    10.4%;
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                                                                   Score 108.5;
Pred. No. 0.36
27; Mismatches
                                                                                 .36;
                                                                                                    DB 7;
                                                                      58;
                                                                      Indels
                                                                                                      Length 860;
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RESULT 4
US-11-069-834-52
(US-11-069-834-52)
(Sequence 52, Application US/11069834)
(Publication No. US20050276811A1)
(PUBLICANT: LIMPORMATION:
(APPLICANT: MOORE JR., FRANCIS D.)
(APPLICANT: HECHTMAN, HERBERT B.)
(APPLICANT: HECHTMAN, HERBERT B.)
(TITLE OF INVENTION: NATURAL IGM ANTIBODIES AND FILE REFERENCE: CRA-002.01)
(CURRENT APPLICATION NUMBER: US/11/069,834)
(CURRENT FILING DATE: 2005-03-01)
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; TYPE: PRT
; ORGANISM: P. :
US-11-196-400-3
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Best Local Similarity
Matches 47; Conserve
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CURRENT FILING DATE: 2005-08-04
PRIOR APPLICATION NUMBER: US/09/742,096
PRIOR FILING DATE: 2000-12-2
PRIOR APPLICATION NUMBER: US 08/973,462
PRIOR FILING DATE: 1998-02-06
PRIOR APPLICATION NUMBER: PCT/FR96/00894
PRIOR FILING DATE: 1996-06-12
PRIOR APPLICATION NUMBER: FR 95/07007
PRIOR FILING DATE: 1996-06-13
PRIOR FILING DATE: 1995-06-13
NUMBER OF SEQ ID NOS: 29
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SEQ ID NO 3
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TITLE OF INVENTION: MALARIAL PRE-ERYTHROCYTIC STAGE POLYPEPTIDE MOLECULES
FILE REFERENCE: 200773USODIV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    1346
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               104 QKSIEKA-----KIGRCETEERTRLAKELSSLRDQREQLKAEVEKYKDCDPQVVEEI 155
                                                                                                                                                                                                                                                                                                                                                                                                    194 TEGIPEDEDYI 204
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  53 SLVDD----GMVDCERIGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQ-----KHASL 103
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EKVIEEEHDITTTLDEVVELKDVEEDKIEK-VSDLKDLEEDILKEVKEIKELESEILEDY 1345
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -AKEAANRWTDNIFAIKSWAKRKFGFEENKIDR 193
                                                                                AND INHIBITORS THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             77;
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US-11-124-368A-214
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TYPE: PRT
ORGANISM: Mus musculus
US-11-069-834-52
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Matches
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APPLICANT: Michele Cargill
APPLICANT: James J. Devlir
APPLICANT: May Luke
                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 214
LENGTH: 2665
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                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Publication No.
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                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Homo sapiens -11-124-368A-214
                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Genetic Polymorphisms Associated with TITLE OF INVENTION: Vascular Diseases, Methods of Detectific Reference: CL001524
CURRENT APPLICATION NUMBER: US/11/124,368A
CURRENT FILING DATE: 2005-05-09
PRIOR APPLICATION NUMBER: US 60/568,845
PRIOR FILING DATE: 2004-05-07
PRIOR APPLICATION NUMBER: US 60/625,936
PRIOR PRIOR PRIOR PRIOR APPLICATION NUMBER: US 60/625,936
PRIOR PRIOR PRIOR DATE: 2004-11-09
NUMBER OF SEQ ID NOS: 21112
SOFTWARE: FASTSEQ for Windows Version 4.0
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PRIOR FILING DATE: 2004-03-01
NUMBER OF SEQ ID NOS: 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver.
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                                                                                                                                                                                                                                                                 Local
                                                                            1794 RGIVSEKTDKLSNMQKDLENSNAKLQEKIQELKANEHQLITLKKDVNETQKKVSEMEQLK 1853
                                                                                                                                                               1737 IDKLRGIVSEKTNEISNMQKDLEHSNDALKAQDLK--IQEELRIAHMHLKE-QQETIDKL
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                                                                                                                                                                                                   1 MSKKKGLSAEEKR--TRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVD--
                                                                                                                                                                                                                                                                 Similarity
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  KQIKDQSLTLSKLEIENLNLAQKLHENLEEMKSVMKERDNLRRVEETLKLERDQLKESLQ 1913
                                          EKAK----IGRCETE-----
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                                                                                                                                                                                                                                                Conservative
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                                                                                                                     DGMVDCER-IGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQKSI 107
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24.7%; Pred. No. 4;
                                                                                                                                                                                                                                            9.6%; Score 100.5;
20.3%; Pred. No. 5.7;
tive 49; Mismatches
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                                    ----ERTRLAKELSSLRDQREQLKAEVE 142
                                                                                                                                                                                                                                                                                    DB 7;
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                                                                                                                                                                                                                                                                                      Length 2665;
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RESULT 7
US-11-069-834-54
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              SEQ ID NO 54
LENGTH: 1976
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                                                                              APPLICANT: CARROLL, MICHAEL C.
APPLICANT: MOORE JR., FRANCIS D.
APPLICANT: HECHTMAN, HERBERT B.
TITLE OF INVENTION: NATURAL IGM ANTIBODIES AND INHIBITORS THEREOF FILE REFERENCE: CRA-002.01
CURRENT APPLICATION NUMBER: US/11/069,834
CURRENT FILING DATE: 2005-03-01
PRIOR APPLICATION NUMBER: 60/588,648
PRIOR APPLICATION NUMBER: 60/549,123
PRIOR PILING DATE: 2004-03-01
RIOR APPLICATION NUMBER: 60/549,123
PRIOR FILING DATE: 2004-03-01
NUMBER: OF SEQ ID NOS: 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/11/124,368A
CURRENT FILING DATE: 2005-05-09
PRIOR APPLICATION NUMBER: US 60/568,845
PRIOR FILING DATE: 2004-05-07
PRIOR PELICATION NUMBER: US 60/625,936
PRIOR APPLICATION NUMBER: US 60/625,936
PRIOR FILING DATE: 2004-11-09
NUMBER OF SEQ ID NOS: 21112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Genetic Polymorphisms Associated with TITLE OF INVENTION: Vascular Diseases, Methods of Detection and Uses Thereof FILE REFERENCE: CL001524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Michele Cargill APPLICANT: James J. Devli
                                                              SOFTWARE: PatentIn Ver.
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TYPE: PRT
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Local Similarity 20.7%;
es 42; Conservation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KKQIKDQSLTLSKLEIENLNLAQKLHENLEEMKSVMKERDNLRRVEETLKLERDQLKESL 1916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RGIVSEKTDKLSNMQKDLENSNAKLQEKVIQELKANEHQLITLKKDVNETQKKVSEMEQL 1856
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Pred. No. 8.8;
47; Mismatches
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; Sequence 994, Application US/10821234; Publication No. US20050255114A1; GENERAL INFORMATION:
; APPLICANT: Labat, Ivan
APPLICANT: Stache-Crain, Birgit; APPLICANT: Andarmani, Susan
; APPLICANT: Tang, Y. Tom
; TITLE OF INVENTION: Methods for Diagnosis and FILE REFERENCE: 821A
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US-11-069-834-48
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US-10-821-234-994
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US-11-069-834-54
                                                     Sequence 48, Application US/11069834
Publication No. US20050276811A1
GENERAL INFORMATION:
APPLICANT: CARROLL, MICHAEL C.
APPLICANT: MOORE JR., FRANCIS D.
APPLICANT: HECHTMAN, HERBERT B.
APPLICANT: HECHTMAN, HERBERT B.
TITLE OF INVENTION: NATURAL IGM ANTIBODIES AND INHIBITORS THEREOF
FILE REFERENCE: CRA-002.01
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Matches
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CURRENT FILING DATE: 2004-04-07
PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 1704
SOFTWARE: pt_SEQ_genes Version 1.0
SEQ ID NO 994
LENGTH: 171
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CURRENT APPLICATION NUMBER: US/11/069,834
CURRENT FILING DATE: 2005-03-01
PRIOR APPLICATION NUMBER: 60/588,648
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             114 RCETEERTRLAKELSSLRDQR----EQLKAEVEKYKDCDPQVVEEIRQANKVAK 163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    878 MERKHQQLLEEKNILAEQLQAETELFAEAEEMRARLAAKKQ-----ELEEILHDLESRVE 932
                                                                                                                                                                                                                                                                                                                                       115 CETEERTRLAKELSSLRDQRE-QLKAEVEKYKDCDPQVVEEIRQANKVAKEAAN 167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ch 9.2%; Score 96; DB 6. I Similarity 20.7%; Pred. No. 0.49; 36; Conservative 38; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EEEER-------NQILQNEKKKMQAHIQDLEEQLDEEEGARQKLQLEKVTAEAKIK 981
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                                                                                                                                                                                                                                                                                                                 KMAEE--KLTHKMEANKENREAOMAAKLERLREKDKH-IEEVRK-NKESKDPAD 166
                                                                                                                                                                                                                                                                                                                                                                                                                                       VDDGMVDCERIGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGR 114
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US-10-821-234-1632
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Best Local :
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                                                                                                                                                                                                                                                                                                             y Match 9.0%; Score 94; DB 6; Length 284;
Local Similarity 24.5%; Pred. No. 1.3;
nes 51; Conservative 30; Mismatches 73; Indels
                       154
                                                                                                                                                                                          59 MVDCERIGISNYYWAFPSKALHAR----KHKLEVLESQLSEGSQKH-------
                                                                                                                                                                                                                                  67
                                                                                                                                                                                                                                                                       2 SKKKGLSAEEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDD----G
                                                                                                    ----ASIQKSIEKAKIGRCETEERTR-LAKELSSLRDQREQLKAEVEKYKDCDPQVVE 153
                   EIRQANKVAKEAANRWTDNIFAIKSWAK 181
                                                                KLVILEGELERSEERAEV----AESRARQLEEELRTMDQALKSLMASEEEYSTKEDKYES 223
                                                                                                                                                                                                                                AQEKLEQAEKKATDABADVASLNRRIQLVEEELDRAQERLATALQKLEBABKAADBSBRG
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------KVIENRAMKDEEKMELQEMQLKEA--KHIAEDSDRKYEEVAR 167

100 126 Gaps

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; TYPE: PRT ; ORGANISM: Mus musculus US-11-069-834-48
CURRENT APPLICATION NUMBER: US/10/821, 234
CURRENT FILING DATE: 2004-04-07
PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
PRIOR FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 1704
SOFTWARE: pt_SEQ_genes Version 1.0
SEQ ID NO 1632
LENGTH: 284
TYPE- DEF
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PRIOR APPLICATION NUMBER: 60/549,123
PRIOR FILING DATE: 2004-03-01
NUMBER OF SEQ ID NOS: 65
SOFTWARE: Patentin Ver. 3.3
SEQ ID NO 48
                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1632, Application US/10821234
Publication No. US20050255114A1
                                                                                                                                                                                                                                 APPLICANT: Labat, Ivan
APPLICANT: Stache-Crain, Birgit
APPLICANT: Andarmani, Susan
APPLICANT: Tang, Y. Tom
APPLICANT: Tang, Y. Methods for Diagnosis and
                                                                                                                                                                                                                    FILE REFERENCE: 821A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 24.1%; Pred. No. 9.6; hes 51; Conservative 37; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            103 LOKSIE----KAKIGRCETBERTRLAKELSSLRDORBOLKAEVEKYKDCDPOVVEEIR- 156
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ----OANKVAKEAANRWIDNIFAIKSWAKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LARALEEAMEQKAELERLNKQFRTEM-EDLMSSKDDVGKSVHELEKSKRALEQQVEEMKT 1539
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EIKLLEEKLKEAETRAE---FAERSVAK 248

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US-10-878-556A-175; Sequence 175, Application No. US20; Publication No. US20; GENERAL INFORMATION:
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                                                                                  ; LENGTH: 1992
; TYPE: PRT
; ORGANISM: Mus musculus
US-11-069-834-58
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CURRENT FILING DATE: 2004-06-28
NUMBER OF SEQ ID NOS: 19
SOFTWARE: PatentIn version 3:1
SEQ ID NO 175
                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/549,123
PRIOR FILING DATE: 2004-03-01
NUMBER OF SEQ ID NOS: 65
SOFTWARE: Patentin Ver. 3.3
SEQ ID NO 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 58, Application US/11069834
Publication No. US20050276811A1
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      Query Match

9.7%; Score 91.5;

Best Local Similarity 24.7%; Pred. No. 20;

Matches 45; Conservative 29; Mismatches
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TYPE: PRT
ORGANISM: Homo sapiens
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: humangp/chrl-aahl5403
DATABASE ENTRY DATE: 2003-04-22
                                                                                                                                                                                                                                                                                                          APPLICANT: CARROLL, MICHAEL C.
APPLICANT: MOORE JR., FRANCIS D.
APPLICANT: HECHYMAN, HERBERT B.
TITLE OF INVENTION: NATURAL IGM ANTIBODIES AND INHIBITORS THEREOF
FILE REFERENCE: CRA-002.01
CURRENT APPLICATION NUMBER: US/11/069,834
CURRENT FILING DATE: 2005-03-01
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TITLE OF INVENTION: HCV regulated protein expression
FILE REFERENCE: 21762
                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/588,648 PRIOR FILING DATE: 2004-07-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      50;
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29;
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Pred. No. 1.3;
3; Mismatches 6
                                            DB 7; Length 1992;
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        61;
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        Indels
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RESULT 14
US-10-878-556A-44
JS-10-878-556A-44
JS-10-878-556A-44
JS-10-878-44
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JS-10
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US-11-069-834-56
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APPLICANT: Hoffmann La-Roche Inc.
TITLE OF INVENTION: HCV regulated protein expression
FILE REFERENCE: 21762
CURRENT APPLICATION NUMBER: US/10/878,556A
CURRENT FILING DATE: 2004-06-28
NUMBER OF SEQ ID NOS: 199
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PRIOR FILING DATE: 2004-03-01
NUMBER OF SEQ ID NOS: 65
SOFTWARE: Patentin Ver. 3.3
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CURRENT APPLICATION NUMBER: US/11/069,834
CURRENT FILING DATE: 2005-03-01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: NATURAL IGM ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Mus
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24.7%; Pred. No. 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29;
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RESULT 15
US-11-669-834-60
US-11-669-834-60
J Sequence 60, Application US/11069834
Publication No. US20050276811A1
J GENERAL INFORMATION:
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-069-834-60
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TYPE: PRT
; ORGANISM: Homo sapiens
; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: sw hum/lama_human
; DATABASE ENTRY DATE: 1986-07-21
US-10-878-556A-44
                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/549,123
PRIOR FILING DATE: 2004-03-01
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn Ver. 3.3
SEQ ID NO 60
SEQ ID NO 60
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                                                                                                                                                                                                                                                                         Query Match 8.7%; Score 91; DB 7; Length 1995; Best Local Similarity 23.9%; Pred. No. 22; Matches 43; Conservative 33; Mismatches 50; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: CARROLL, MICHAEL C.
APPLICANT: MOORE JR., FRANCIS D.
APPLICANT: HECHTMAN, HERBERT B.
TITLE OF INVENTION: NATURAL IGM ANTIBODIES AND INHIBITORS THEREOF
FILE REFERENCE: CRA-002.01
CURRENT APPLICATION NUMBER: US/11/069,834
CURRENT FILING DATE: 2005-03-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/588,648 PRIOR FILING DATE: 2004-07-16
1500 ARALSITRALBEEQEAREBLERONRALRAELEALISSKODVGKSVHELERACRVAEQAAN 1559
                                                                                          1459 LVST------LEKKORKFDOLL-----AEEKAAVLRAVEERE--RAEAEGRERE
                                                                                                                                                                                 1414 AAREAEALTQRLAEKTETVDRL------ERG-----RRRLQQELDDATMDLEQQRQ 1458
                                               124 AKELS-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  154 EKRTLEGELHDLRGQVAKLEAALGEAKKQLQDEMLRRVDAEN------RLQTMKEELD 205
                                                                                                                                    65 -IGTSNYYWAFPSKALHARKHKLEVLESQLSEGSQKHASLQKSIEKAKIGRCETEERTRL 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88 ----VLESQLSEGSQKHASLQKSIEKAKIGRCETEERTRLAKELSSLR----DQREQLKA 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             47 VKEVLQSLVDD--GMVD------CERIGTSNYYWAFPSKALHARKHKLE
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                                                                                                                                                                                                                              8 SAEEKRTRMMEIFSETKDVFQLKDLEKIAPKEKGITAMSVKEVLQSLVDDGMVDCER--- 64
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                            ----SLRDQREQLKAEVE---KYKDCDPQVVEBIRQANKVAKEAAN 167
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Search completed: February 2, 2006, 12:56:28

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